

Machine Learning Lab

Assignment 3

Name – SHAURYA PRAKASH SHAH

Roll - 001811001025

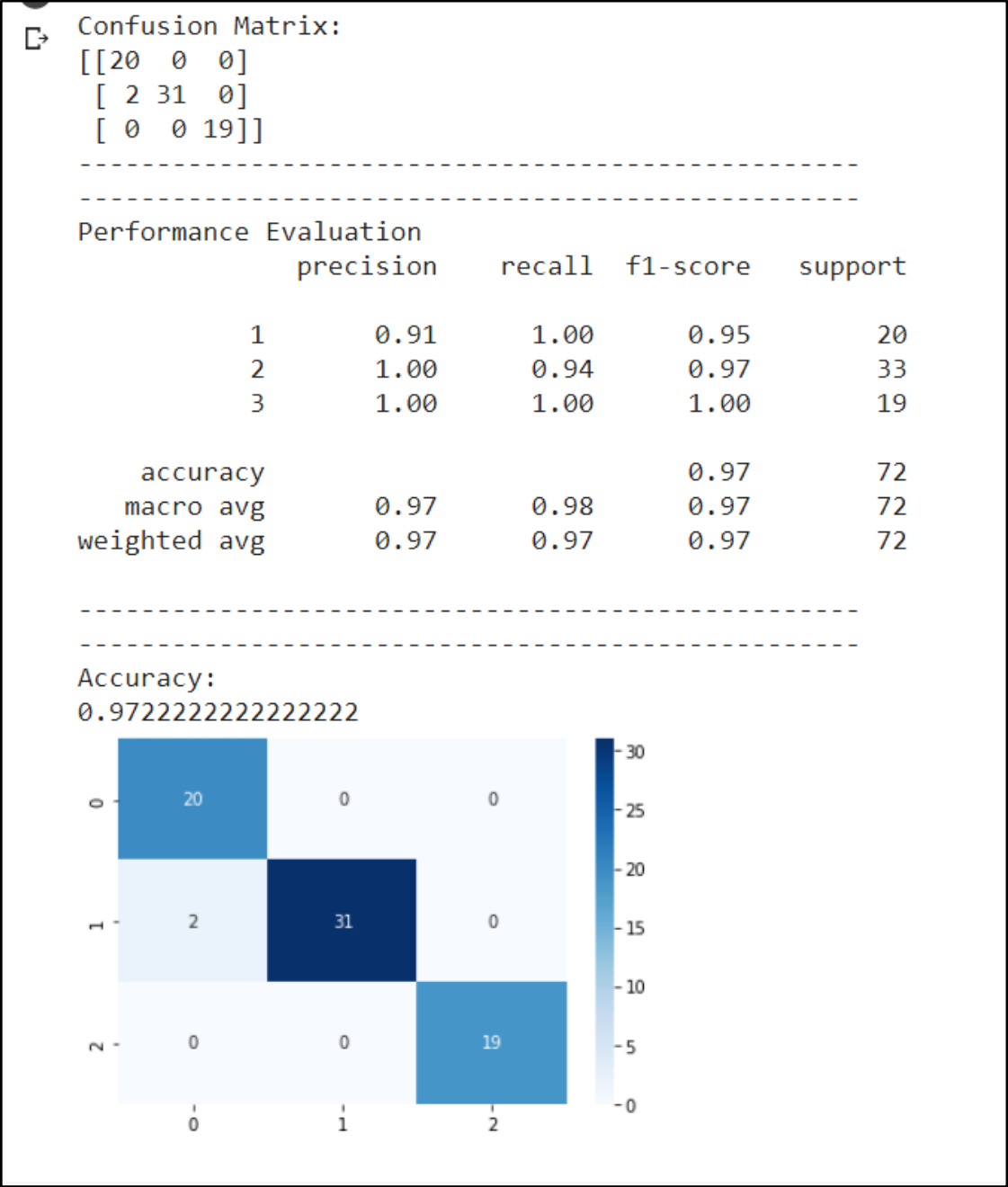
Semester - 7

Year - 4

Department - Information Technology

1) Wine Dataset

1.1) GaussianHMM Without Tuning



1.2) GaussianHMM With Tuning

Confusion Matrix:

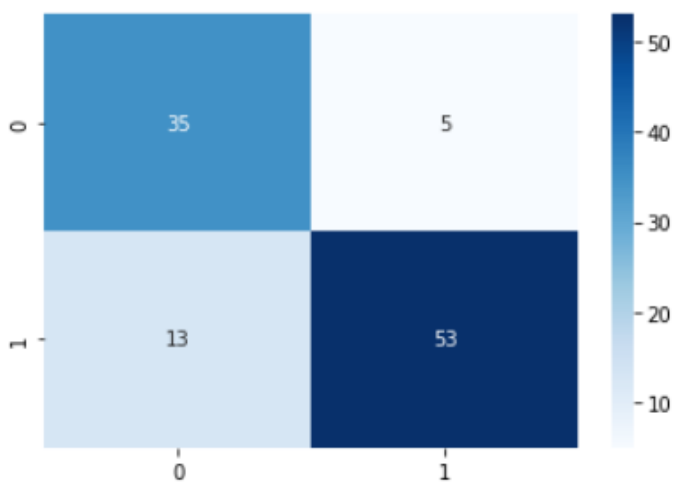
```
[[35  5]
 [13 53]]
```

Performance Evaluation

	precision	recall	f1-score	support
b	0.73	0.88	0.80	40
g	0.91	0.80	0.85	66
accuracy			0.83	106
macro avg	0.82	0.84	0.83	106
weighted avg	0.84	0.83	0.83	106

Accuracy:

0.8301886792452831



1.3) GMMHMM Without Tuning

Confusion Matrix:

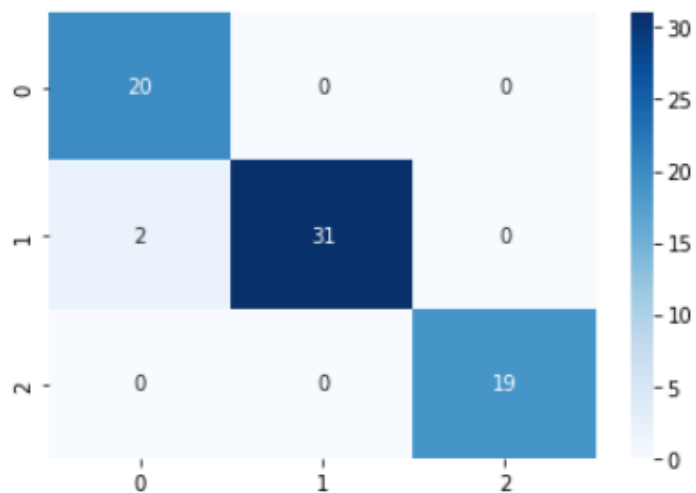
```
[[20  0  0]
 [ 2 31  0]
 [ 0  0 19]]
```

Performance Evaluation

	precision	recall	f1-score	support
1	0.91	1.00	0.95	20
2	1.00	0.94	0.97	33
3	1.00	1.00	1.00	19
accuracy			0.97	72
macro avg	0.97	0.98	0.97	72
weighted avg	0.97	0.97	0.97	72

Accuracy:

0.9722222222222222



1.4) GMMHMM With Tuning

➔ Confusion Matrix:

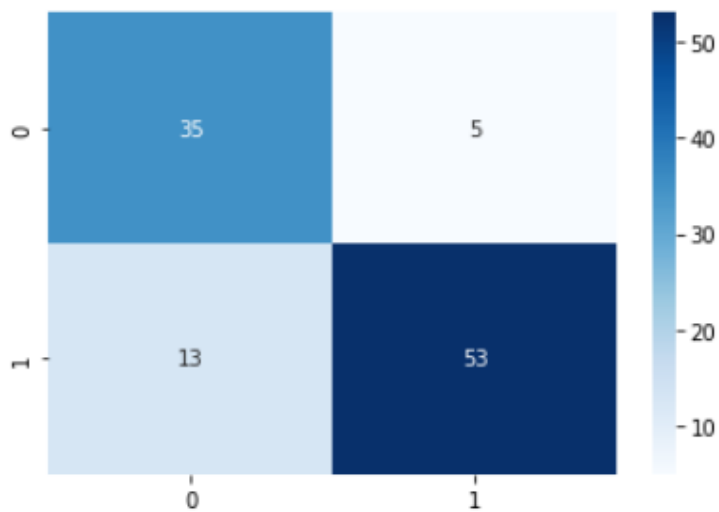
```
[[35  5]
 [13 53]]
```


Performance Evaluation

	precision	recall	f1-score	support
b	0.73	0.88	0.80	40
g	0.91	0.80	0.85	66
accuracy			0.83	106
macro avg	0.82	0.84	0.83	106
weighted avg	0.84	0.83	0.83	106

Accuracy:

0.8301886792452831



1.5) MultinomialHMM Without Tuning

↳ Confusion Matrix:

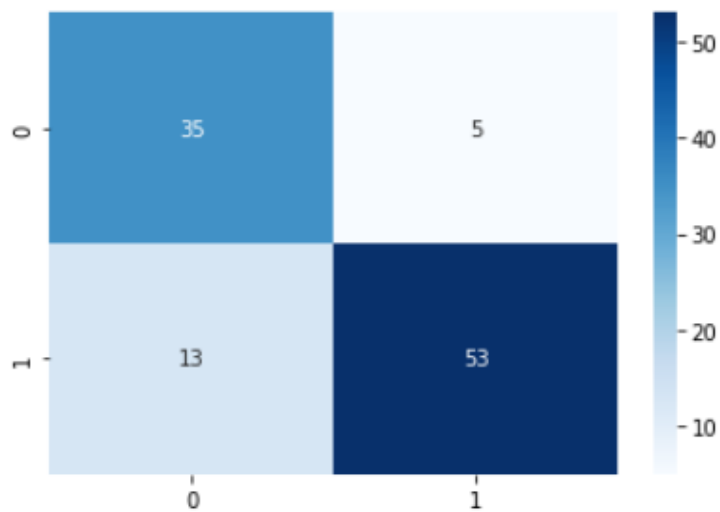
```
[[35  5]
 [13 53]]
```


Performance Evaluation

	precision	recall	f1-score	support
b	0.73	0.88	0.80	40
g	0.91	0.80	0.85	66
accuracy			0.83	106
macro avg	0.82	0.84	0.83	106
weighted avg	0.84	0.83	0.83	106

Accuracy:

0.8301886792452831



1.6) MultinomialHMM Without Tuning

➞ Confusion Matrix:

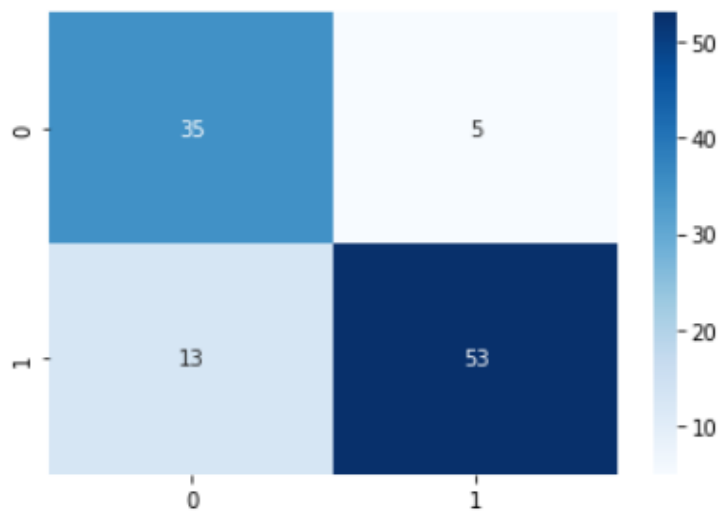
```
[[35  5]
 [13 53]]
```


Performance Evaluation

	precision	recall	f1-score	support
b	0.73	0.88	0.80	40
g	0.91	0.80	0.85	66
accuracy			0.83	106
macro avg	0.82	0.84	0.83	106
weighted avg	0.84	0.83	0.83	106

Accuracy:

0.8301886792452831



2.1) GaussianHMM Without Tuning

➔ Confusion Matrix:

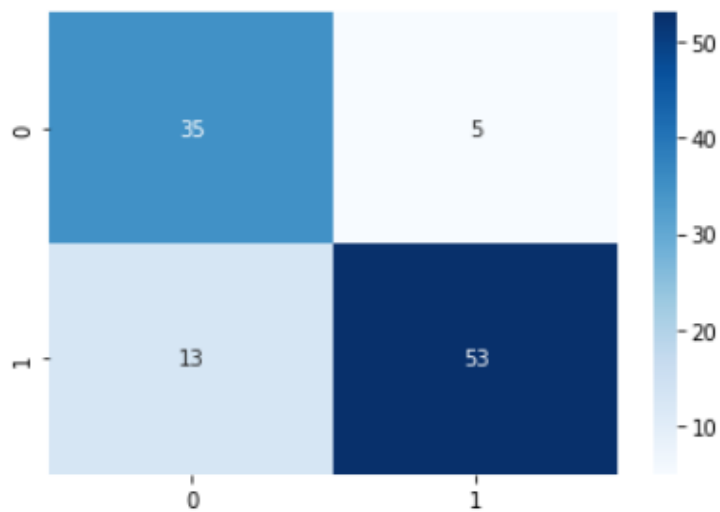
```
[[35  5]
 [13 53]]
```


Performance Evaluation

	precision	recall	f1-score	support
b	0.73	0.88	0.80	40
g	0.91	0.80	0.85	66
accuracy			0.83	106
macro avg	0.82	0.84	0.83	106
weighted avg	0.84	0.83	0.83	106

Accuracy:

0.8301886792452831



2.2) GaussianHMM With Tuning

➤ Confusion Matrix:

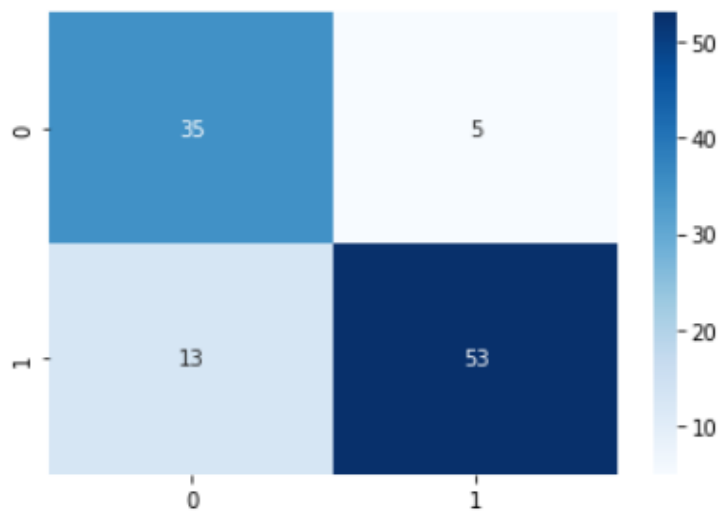
```
[[35  5]
 [13 53]]
```


Performance Evaluation

	precision	recall	f1-score	support
b	0.73	0.88	0.80	40
g	0.91	0.80	0.85	66
accuracy			0.83	106
macro avg	0.82	0.84	0.83	106
weighted avg	0.84	0.83	0.83	106

Accuracy:

0.8301886792452831



2.3) GMMHMM Without Tuning

➤ Confusion Matrix:

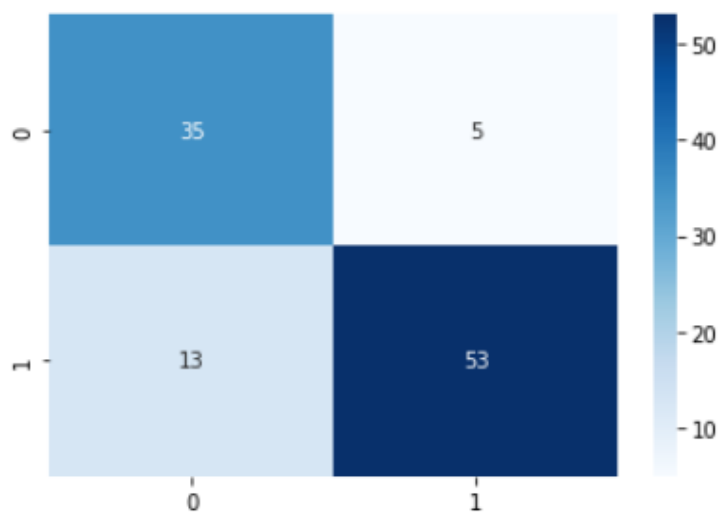
```
[[35  5]
 [13 53]]
```


Performance Evaluation

	precision	recall	f1-score	support
b	0.73	0.88	0.80	40
g	0.91	0.80	0.85	66
accuracy			0.83	106
macro avg	0.82	0.84	0.83	106
weighted avg	0.84	0.83	0.83	106

Accuracy:

0.8301886792452831



2.4) GMMHMM With Tuning

➤ Confusion Matrix:

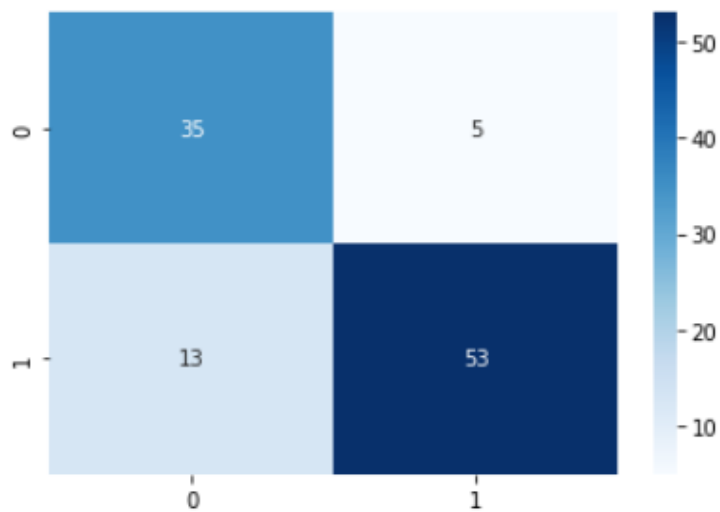
```
[[35  5]
 [13 53]]
```


Performance Evaluation

	precision	recall	f1-score	support
b	0.73	0.88	0.80	40
g	0.91	0.80	0.85	66
accuracy			0.83	106
macro avg	0.82	0.84	0.83	106
weighted avg	0.84	0.83	0.83	106

Accuracy:

0.8301886792452831



2.5) MultinomialHMM Without Tuning

➤ Confusion Matrix:

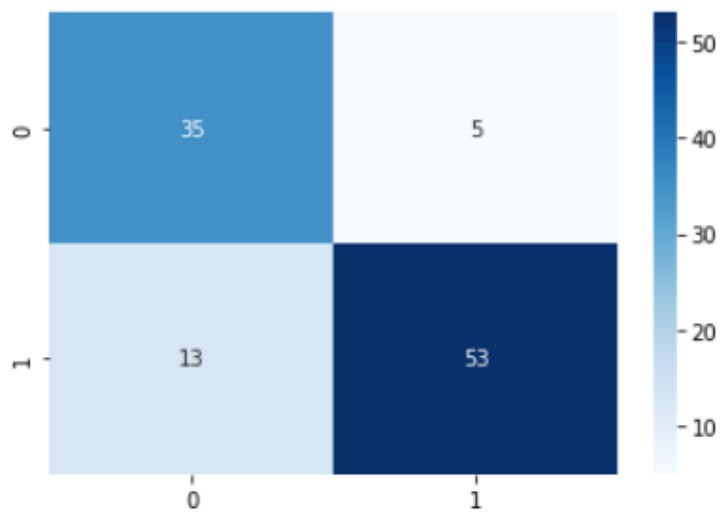
```
[[35  5]
 [13 53]]
```


Performance Evaluation

	precision	recall	f1-score	support
b	0.73	0.88	0.80	40
g	0.91	0.80	0.85	66
accuracy			0.83	106
macro avg	0.82	0.84	0.83	106
weighted avg	0.84	0.83	0.83	106

Accuracy:

0.8301886792452831



2.6) MultinomialHMM Without Tuning

Confusion Matrix:

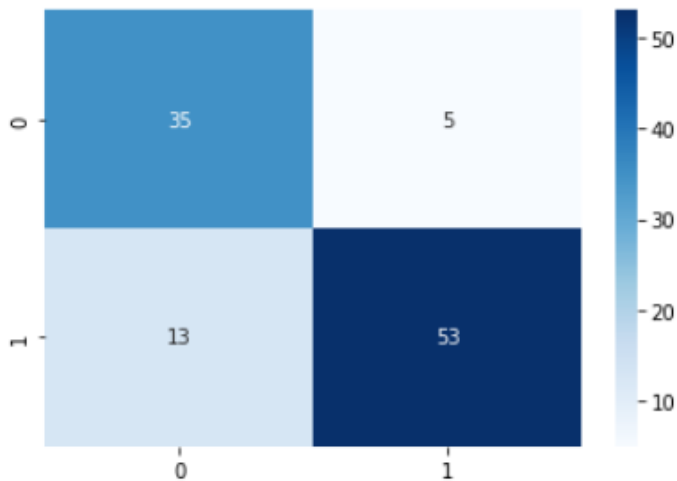
```
[[35  5]
 [13 53]]
```

Performance Evaluation

	precision	recall	f1-score	support
b	0.73	0.88	0.80	40
g	0.91	0.80	0.85	66
accuracy			0.83	106
macro avg	0.82	0.84	0.83	106
weighted avg	0.84	0.83	0.83	106

Accuracy:

0.8301886792452831



3) Breast Cancer Dataset

3.1) GaussianHMM Without Tuning

Confusion Matrix:

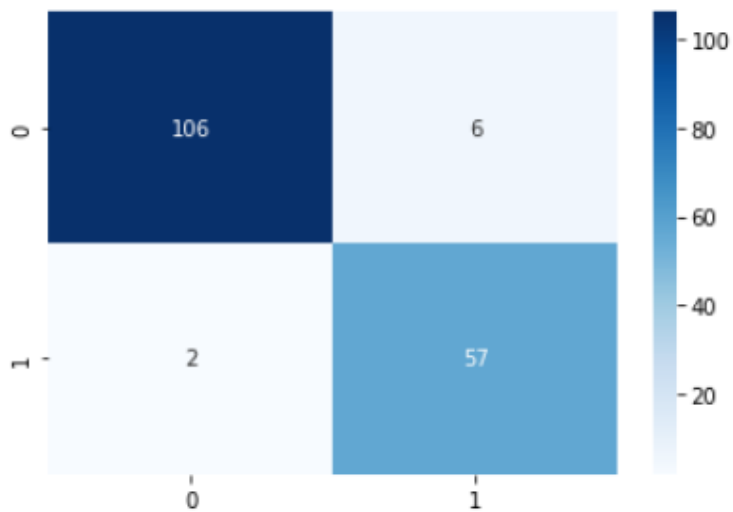
```
[[106  6]
 [  2 57]]
```

Performance Evaluation

	precision	recall	f1-score	support
B	0.98	0.95	0.96	112
M	0.90	0.97	0.93	59
accuracy			0.95	171
macro avg	0.94	0.96	0.95	171
weighted avg	0.96	0.95	0.95	171

Accuracy:

0.9532163742690059



3.2) GaussianHMM With Tuning

Confusion Matrix:

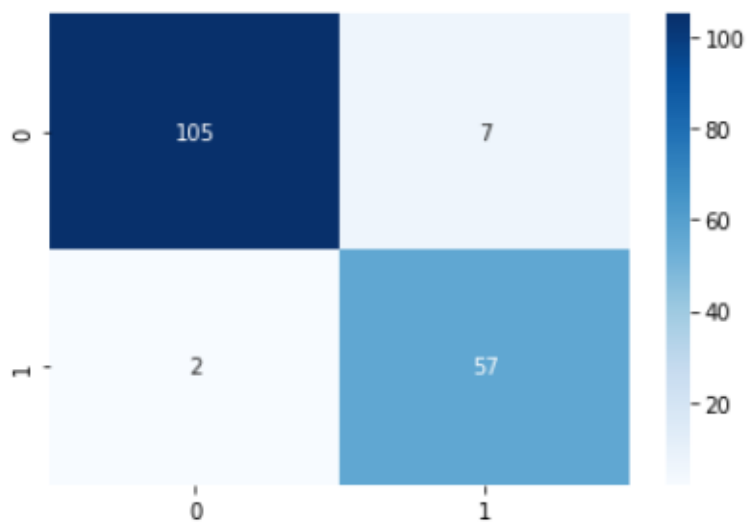
```
[[105  7]
 [  2 57]]
```

Performance Evaluation

	precision	recall	f1-score	support
B	0.98	0.94	0.96	112
M	0.89	0.97	0.93	59
accuracy			0.95	171
macro avg	0.94	0.95	0.94	171
weighted avg	0.95	0.95	0.95	171

Accuracy:

0.9473684210526315



3.3) GMMHMM Without Tuning

↳ Confusion Matrix:

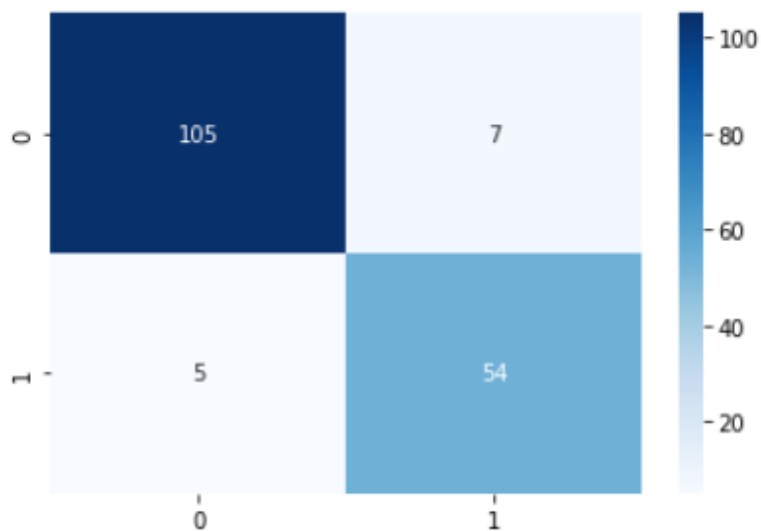
```
[[105  7]
 [ 5 54]]
```


Performance Evaluation

	precision	recall	f1-score	support
B	0.95	0.94	0.95	112
M	0.89	0.92	0.90	59
accuracy			0.93	171
macro avg	0.92	0.93	0.92	171
weighted avg	0.93	0.93	0.93	171

Accuracy:

0.9298245614035088



3.4) GMMHMM With Tuning

↳ Confusion Matrix:

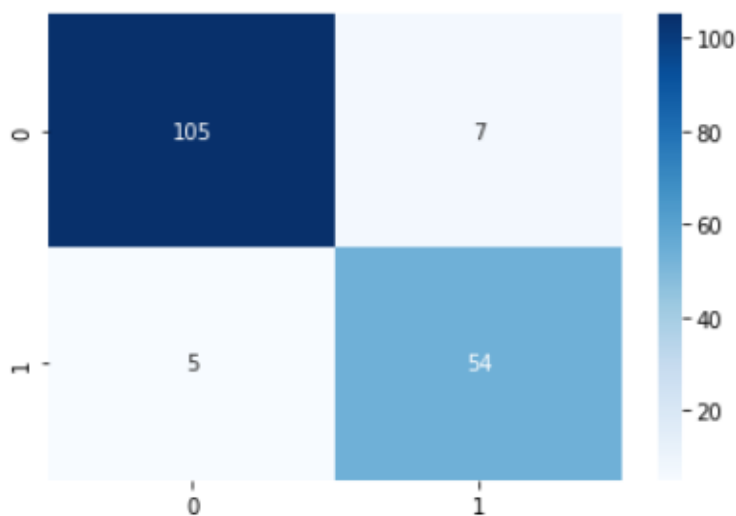
```
[[105  7]
 [ 5 54]]
```

Performance Evaluation

	precision	recall	f1-score	support
B	0.95	0.94	0.95	112
M	0.89	0.92	0.90	59
accuracy			0.93	171
macro avg	0.92	0.93	0.92	171
weighted avg	0.93	0.93	0.93	171

Accuracy:

0.9298245614035088



3.5) MultinomialHMM Without Tuning

Confusion Matrix:

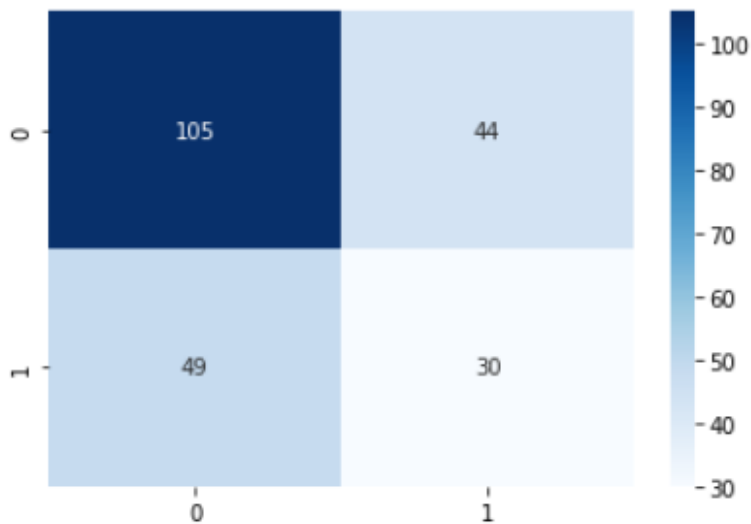
```
[[105  44]
 [ 49  30]]
```

Performance Evaluation

	precision	recall	f1-score	support
B	0.68	0.70	0.69	149
M	0.41	0.38	0.39	79
accuracy			0.59	228
macro avg	0.54	0.54	0.54	228
weighted avg	0.59	0.59	0.59	228

Accuracy:

0.5921052631578947



3.6) MultinomialHMM Without Tuning

Confusion Matrix:

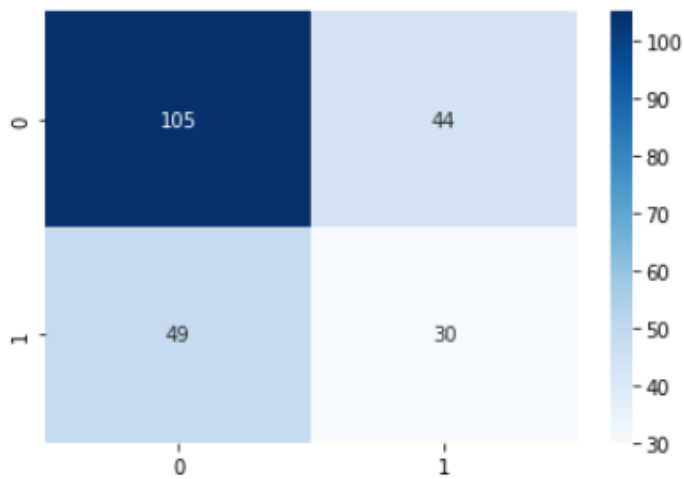
```
[[105  44]
 [ 49  30]]
```

Performance Evaluation

	precision	recall	f1-score	support
B	0.68	0.70	0.69	149
M	0.41	0.38	0.39	79
accuracy			0.59	228
macro avg	0.54	0.54	0.54	228
weighted avg	0.59	0.59	0.59	228

Accuracy:

0.5921052631578947

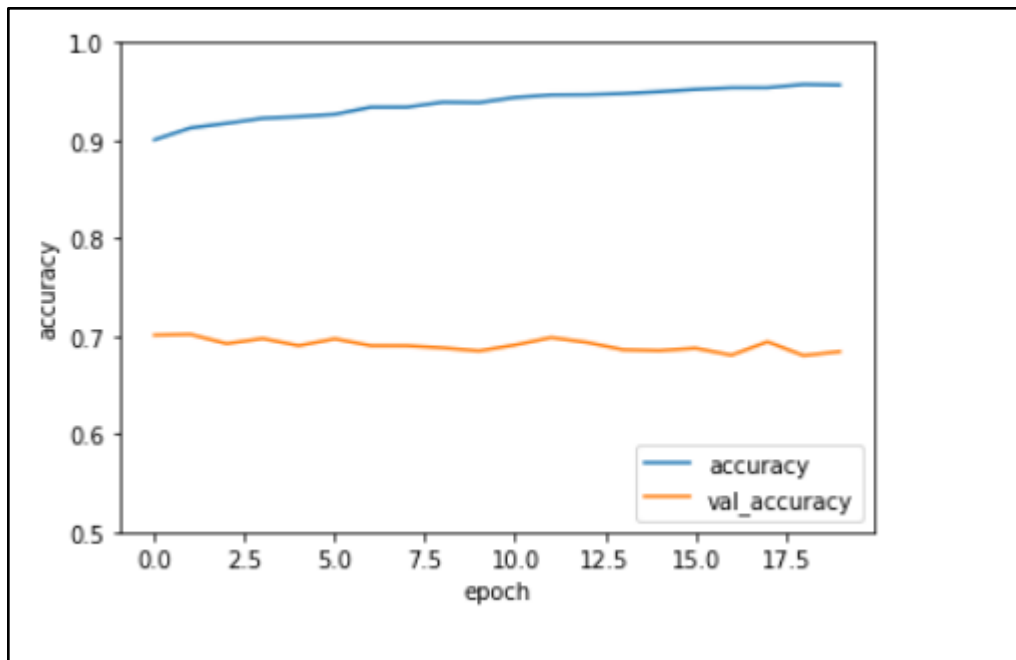


1) CIFAR-10

Model: "sequential_2"

Layer (type)	Output Shape	Param #
conv2d_6 (Conv2D)	(None, 30, 30, 32)	896
max_pooling2d_4 (MaxPooling2D)	(None, 15, 15, 32)	0
conv2d_7 (Conv2D)	(None, 13, 13, 64)	18496
max_pooling2d_5 (MaxPooling2D)	(None, 6, 6, 64)	0
conv2d_8 (Conv2D)	(None, 4, 4, 64)	36928
flatten (Flatten)	(None, 1024)	0
dense (Dense)	(None, 64)	65600
dense_1 (Dense)	(None, 10)	650
Total params: 122,570		
Trainable params: 122,570		
Non-trainable params: 0		

```
Epoch 11/20
1563/1563 [=====] - 69s 44ms/step - loss: 0.1561 - accuracy: 0.9437 - val_loss: 1.8716 - val_accuracy: 0.6910
Epoch 12/20
1563/1563 [=====] - 69s 44ms/step - loss: 0.1498 - accuracy: 0.9463 - val_loss: 1.9775 - val_accuracy: 0.6986
Epoch 13/20
1563/1563 [=====] - 69s 44ms/step - loss: 0.1524 - accuracy: 0.9466 - val_loss: 2.0503 - val_accuracy: 0.6936
Epoch 14/20
1563/1563 [=====] - 69s 44ms/step - loss: 0.1490 - accuracy: 0.9477 - val_loss: 2.0715 - val_accuracy: 0.6861
Epoch 15/20
1563/1563 [=====] - 69s 44ms/step - loss: 0.1429 - accuracy: 0.9497 - val_loss: 2.1616 - val_accuracy: 0.6852
Epoch 16/20
1563/1563 [=====] - 69s 44ms/step - loss: 0.1356 - accuracy: 0.9520 - val_loss: 2.2363 - val_accuracy: 0.6877
Epoch 17/20
1563/1563 [=====] - 69s 44ms/step - loss: 0.1327 - accuracy: 0.9537 - val_loss: 2.2239 - val_accuracy: 0.6806
Epoch 18/20
1563/1563 [=====] - 69s 44ms/step - loss: 0.1348 - accuracy: 0.9538 - val_loss: 2.2102 - val_accuracy: 0.6943
Epoch 19/20
1563/1563 [=====] - 69s 44ms/step - loss: 0.1250 - accuracy: 0.9571 - val_loss: 2.3900 - val_accuracy: 0.6802
Epoch 20/20
1563/1563 [=====] - 69s 44ms/step - loss: 0.1245 - accuracy: 0.9565 - val_loss: 2.3173 - val_accuracy: 0.6842
```



2) MNIST

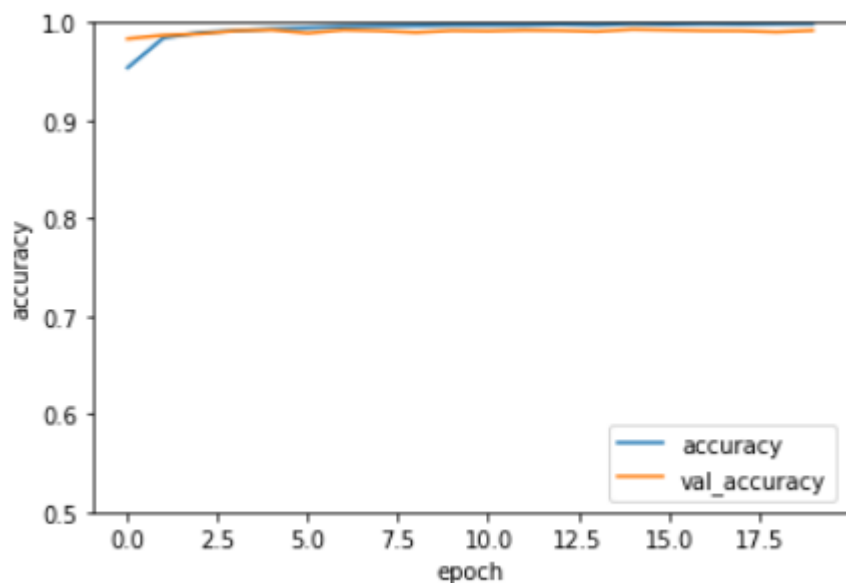
Model: "sequential_8"

Layer (type)	Output Shape	Param #
conv2d_18 (Conv2D)	(None, 26, 26, 32)	320
max_pooling2d_10 (MaxPooling)	(None, 13, 13, 32)	0
conv2d_19 (Conv2D)	(None, 11, 11, 64)	18496
max_pooling2d_11 (MaxPooling)	(None, 5, 5, 64)	0
conv2d_20 (Conv2D)	(None, 3, 3, 64)	36928
flatten_3 (Flatten)	(None, 576)	0
dense_6 (Dense)	(None, 64)	36928
dense_7 (Dense)	(None, 10)	650
Total params: 93,322		
Trainable params: 93,322		
Non-trainable params: 0		

```

1875/1875 [=====] - 57s 30ms/step - loss: 0.0079 - accuracy: 0.9973 - val_loss: 0.0319 - val_accuracy: 0.9913
Epoch 12/20
1875/1875 [=====] - 57s 31ms/step - loss: 0.0084 - accuracy: 0.9973 - val_loss: 0.0329 - val_accuracy: 0.9921
Epoch 13/20
1875/1875 [=====] - 57s 31ms/step - loss: 0.0067 - accuracy: 0.9980 - val_loss: 0.0343 - val_accuracy: 0.9918
Epoch 14/20
1875/1875 [=====] - 58s 31ms/step - loss: 0.0078 - accuracy: 0.9973 - val_loss: 0.0390 - val_accuracy: 0.9908
Epoch 15/20
1875/1875 [=====] - 58s 31ms/step - loss: 0.0048 - accuracy: 0.9985 - val_loss: 0.0374 - val_accuracy: 0.9930
Epoch 16/20
1875/1875 [=====] - 58s 31ms/step - loss: 0.0069 - accuracy: 0.9980 - val_loss: 0.0336 - val_accuracy: 0.9923
Epoch 17/20
1875/1875 [=====] - 57s 31ms/step - loss: 0.0049 - accuracy: 0.9985 - val_loss: 0.0430 - val_accuracy: 0.9916
Epoch 18/20
1875/1875 [=====] - 57s 31ms/step - loss: 0.0053 - accuracy: 0.9982 - val_loss: 0.0397 - val_accuracy: 0.9915
Epoch 19/20
1875/1875 [=====] - 58s 31ms/step - loss: 0.0048 - accuracy: 0.9986 - val_loss: 0.0540 - val_accuracy: 0.9903
Epoch 20/20
1875/1875 [=====] - 58s 31ms/step - loss: 0.0043 - accuracy: 0.9987 - val_loss: 0.0419 - val_accuracy: 0.9919

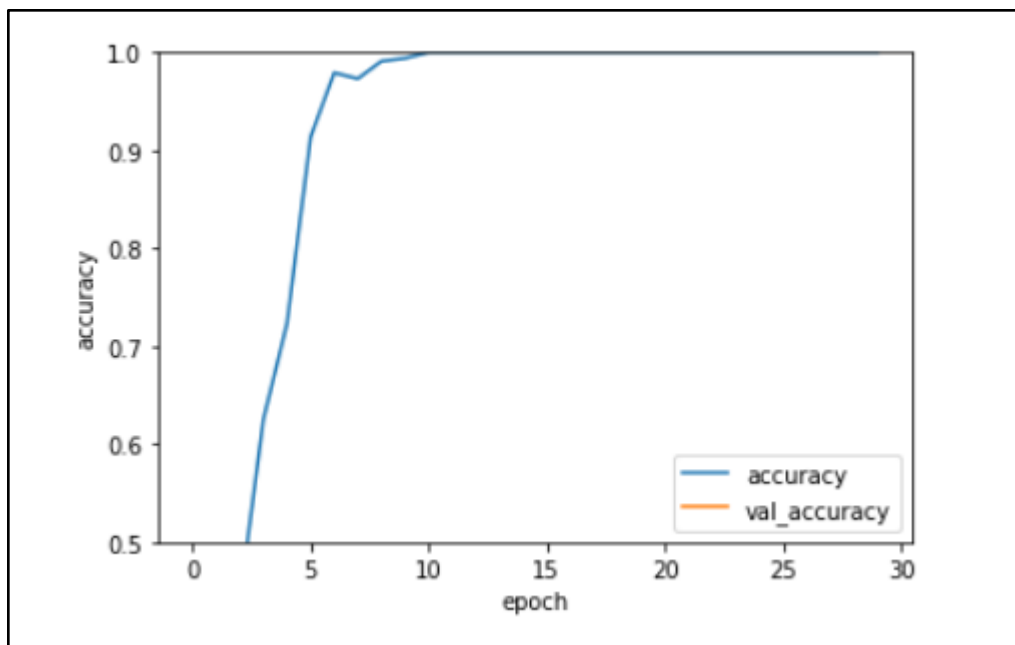
```



3) SAVEE

Model: "sequential_3"		
Layer (type)	Output Shape	Param #
=====		
conv2d_9 (Conv2D)	(None, 155, 318, 32)	320
<hr/>		
max_pooling2d_6 (MaxPooling2)	(None, 77, 159, 32)	0
<hr/>		
conv2d_10 (Conv2D)	(None, 75, 157, 64)	18496
<hr/>		
max_pooling2d_7 (MaxPooling2)	(None, 37, 78, 64)	0
<hr/>		
conv2d_11 (Conv2D)	(None, 35, 76, 64)	36928
<hr/>		
flatten_3 (Flatten)	(None, 170240)	0
<hr/>		
dense_6 (Dense)	(None, 64)	10895424
<hr/>		
dense_7 (Dense)	(None, 10)	650
<hr/>		
Total params: 10,951,818		
Trainable params: 10,951,818		
Non-trainable params: 0		
<hr/>		

11/11	[=====]	- 27s	2s/step	- loss: 2.2025e-05	- accuracy: 1.0000	- val_loss: 7.0087	- val_accuracy: 0.3056
Epoch 25/30							
11/11	[=====]	- 27s	2s/step	- loss: 1.9328e-05	- accuracy: 1.0000	- val_loss: 7.0391	- val_accuracy: 0.2986
Epoch 26/30							
11/11	[=====]	- 27s	2s/step	- loss: 1.7196e-05	- accuracy: 1.0000	- val_loss: 7.0967	- val_accuracy: 0.2986
Epoch 27/30							
11/11	[=====]	- 27s	2s/step	- loss: 1.5431e-05	- accuracy: 1.0000	- val_loss: 7.1239	- val_accuracy: 0.3056
Epoch 28/30							
11/11	[=====]	- 27s	2s/step	- loss: 1.3852e-05	- accuracy: 1.0000	- val_loss: 7.1493	- val_accuracy: 0.2986
Epoch 29/30							
11/11	[=====]	- 27s	2s/step	- loss: 1.2641e-05	- accuracy: 1.0000	- val_loss: 7.2041	- val_accuracy: 0.2986
Epoch 30/30							
11/11	[=====]	- 27s	2s/step	- loss: 1.1668e-05	- accuracy: 1.0000	- val_loss: 7.2112	- val_accuracy: 0.2986



4) EmoDB

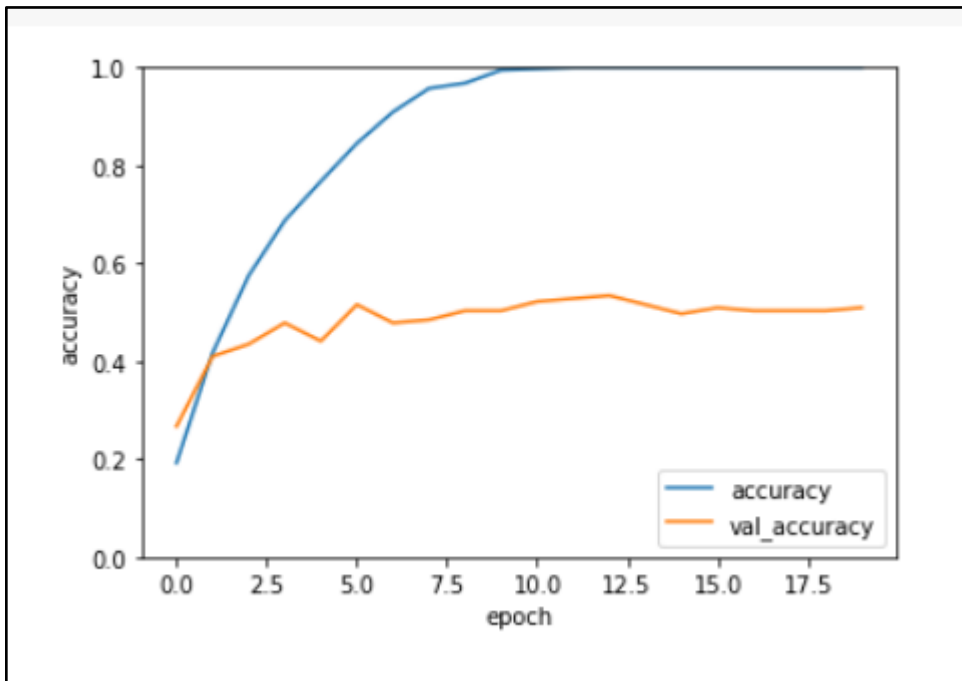
Model: "sequential_4"

Layer (type)	Output Shape	Param #
=====		
conv2d_12 (Conv2D)	(None, 155, 318, 32)	320
max_pooling2d_8 (MaxPooling2D)	(None, 77, 159, 32)	0
conv2d_13 (Conv2D)	(None, 75, 157, 64)	18496
max_pooling2d_9 (MaxPooling2D)	(None, 37, 78, 64)	0
conv2d_14 (Conv2D)	(None, 35, 76, 64)	36928
flatten_4 (Flatten)	(None, 170240)	0
dense_8 (Dense)	(None, 64)	10895424
dense_9 (Dense)	(None, 10)	650
=====		
Total params: 10,951,818		
Trainable params: 10,951,818		
Non-trainable params: 0		


```

Epoch 14/20
12/12 [=====] - 30s 2s/step - loss: 0.0012 - accuracy: 1.0000 - val_loss: 3.9037 - val_accuracy: 0.5155
Epoch 15/20
12/12 [=====] - 30s 2s/step - loss: 7.0827e-04 - accuracy: 1.0000 - val_loss: 4.0446 - val_accuracy: 0.4969
Epoch 16/20
12/12 [=====] - 30s 2s/step - loss: 4.9740e-04 - accuracy: 1.0000 - val_loss: 4.1150 - val_accuracy: 0.5093
Epoch 17/20
12/12 [=====] - 30s 3s/step - loss: 3.8747e-04 - accuracy: 1.0000 - val_loss: 4.1542 - val_accuracy: 0.5031
Epoch 18/20
12/12 [=====] - 30s 2s/step - loss: 3.0542e-04 - accuracy: 1.0000 - val_loss: 4.2023 - val_accuracy: 0.5031
Epoch 19/20
12/12 [=====] - 31s 3s/step - loss: 2.5256e-04 - accuracy: 1.0000 - val_loss: 4.2239 - val_accuracy: 0.5031
Epoch 20/20
12/12 [=====] - 30s 2s/step - loss: 2.1154e-04 - accuracy: 1.0000 - val_loss: 4.2753 - val_accuracy: 0.5093

```



1) VGG-16

1.1) CIFAR-10

```
Epoch 1/5
63/63 [=====] - 97s 855ms/step - loss: nan - accuracy: 0.0985
Epoch 2/5
63/63 [=====] - 47s 743ms/step - loss: nan - accuracy: 0.1010
Epoch 3/5
63/63 [=====] - 47s 745ms/step - loss: nan - accuracy: 0.1010
Epoch 4/5
63/63 [=====] - 47s 744ms/step - loss: nan - accuracy: 0.1010
Epoch 5/5
63/63 [=====] - 47s 744ms/step - loss: nan - accuracy: 0.1010
```

```
[ ] model.evaluate(X_test_resized, y_test)
```

```
63/63 [=====] - 14s 225ms/step - loss: nan - accuracy: 0.0980
[nan, 0.0979999974370027]
```

1.2) MNIST

```
Epoch 1/5
63/63 [=====] - 71s 877ms/step - loss: 3.6193 - accuracy: 0.0885
Epoch 2/5
63/63 [=====] - 48s 764ms/step - loss: 2.5311 - accuracy: 0.0990
Epoch 3/5
63/63 [=====] - 48s 763ms/step - loss: 2.5246 - accuracy: 0.1105
Epoch 4/5
63/63 [=====] - 48s 763ms/step - loss: 2.5036 - accuracy: 0.1040
Epoch 5/5
63/63 [=====] - 48s 763ms/step - loss: 2.4806 - accuracy: 0.0965
```

```
model.evaluate(X_test_new, y_test)
```

```
63/63 [=====] - 15s 233ms/step - loss: 2.6352 - accuracy: 0.1095
[2.6351511478424072, 0.10949999839067459]
```

1.3) SAVEE

```
8/8 [=====] - 6s 708ms/step - loss: nan - accuracy: 0.1208
Epoch 45/50
8/8 [=====] - 6s 706ms/step - loss: nan - accuracy: 0.1208
Epoch 46/50
8/8 [=====] - 6s 705ms/step - loss: nan - accuracy: 0.1208
Epoch 47/50
8/8 [=====] - 6s 706ms/step - loss: nan - accuracy: 0.1208
Epoch 48/50
8/8 [=====] - 6s 709ms/step - loss: nan - accuracy: 0.1208
Epoch 49/50
8/8 [=====] - 6s 706ms/step - loss: nan - accuracy: 0.1208
Epoch 50/50
8/8 [=====] - 6s 706ms/step - loss: nan - accuracy: 0.1208
```

```
model.evaluate(X_test_resized, y_test)
```

```
8/8 [=====] - 2s 215ms/step - loss: nan - accuracy: 0.1292
[nan, 0.12916666269302368]
```

1.4) EmoDB

```
9/9 [=====] - 6s 711ms/step - loss: nan - accuracy: 0.2247
Epoch 14/20
9/9 [=====] - 6s 712ms/step - loss: nan - accuracy: 0.2247
Epoch 15/20
9/9 [=====] - 6s 713ms/step - loss: nan - accuracy: 0.2247
Epoch 16/20
9/9 [=====] - 6s 712ms/step - loss: nan - accuracy: 0.2247
Epoch 17/20
9/9 [=====] - 6s 712ms/step - loss: nan - accuracy: 0.2247
Epoch 18/20
9/9 [=====] - 6s 711ms/step - loss: nan - accuracy: 0.2247
Epoch 19/20
9/9 [=====] - 6s 712ms/step - loss: nan - accuracy: 0.2247
Epoch 20/20
9/9 [=====] - 6s 712ms/step - loss: nan - accuracy: 0.2247
```

```
model.evaluate(X_test_resized, y_test)
```

```
9/9 [=====] - 6s 718ms/step - loss: nan - accuracy: 0.2500
[nan, 0.25]
```

The entire model can be broken down into 5 blocks, where each block contains 3 convolution and 1 max-pooling layers.

Looking at the complexity of the model and the limitations of google colab, I have reduced the input size for the model,i.e., i have taken 2000 training data points and 2000 testing data points.

2) ResNet-50

2.1) CIFAR-10

```
Downloading data from https://github.com/fchollet/deep-learning-models/releases/download/v0.2/resnet50\_102858752/102853048 [=====] - 1s 0us/step
102866944/102853048 [=====] - 1s 0us/step
Epoch 1/5
63/63 [=====] - 81s 703ms/step - loss: 2.9229 - accuracy: 0.0975
Epoch 2/5
63/63 [=====] - 42s 673ms/step - loss: 2.4506 - accuracy: 0.1040
Epoch 3/5
63/63 [=====] - 42s 673ms/step - loss: 2.2995 - accuracy: 0.1755
Epoch 4/5
63/63 [=====] - 42s 672ms/step - loss: 2.1401 - accuracy: 0.2325
Epoch 5/5
63/63 [=====] - 42s 672ms/step - loss: 2.0272 - accuracy: 0.2715

model.evaluate(X_test_resized, y_test)

63/63 [=====] - 15s 217ms/step - loss: 16.8393 - accuracy: 0.0000e+00
[16.839269638061523, 0.0]
```

2.2) MNIST

```

Epoch 6/10
63/63 [=====] - 44s 705ms/step - loss: 0.0647 - accuracy: 0.9825
Epoch 7/10
63/63 [=====] - 44s 705ms/step - loss: 0.0655 - accuracy: 0.9815
Epoch 8/10
63/63 [=====] - 44s 705ms/step - loss: 0.0429 - accuracy: 0.9890
Epoch 9/10
63/63 [=====] - 44s 705ms/step - loss: 0.0272 - accuracy: 0.9950
Epoch 10/10
63/63 [=====] - 44s 704ms/step - loss: 0.0121 - accuracy: 0.9985

```

```

model.evaluate(X_test_new, y_test)

```

```

63/63 [=====] - 14s 227ms/step - loss: 13.1569 - accuracy: 0.0000e+00
[13.156914710998535, 0.0]

```

2.3) SAVEE

```

Epoch 4/10
8/8 [=====] - 5s 669ms/step - loss: 1.0197 - accuracy: 0.6833
Epoch 5/10
8/8 [=====] - 5s 667ms/step - loss: 0.5054 - accuracy: 0.9167
Epoch 6/10
8/8 [=====] - 5s 669ms/step - loss: 0.2390 - accuracy: 0.9833
Epoch 7/10
8/8 [=====] - 5s 671ms/step - loss: 0.0966 - accuracy: 1.0000
Epoch 8/10
8/8 [=====] - 5s 668ms/step - loss: 0.0691 - accuracy: 1.0000
Epoch 9/10
8/8 [=====] - 5s 669ms/step - loss: 0.0550 - accuracy: 1.0000
Epoch 10/10
8/8 [=====] - 5s 666ms/step - loss: 0.0281 - accuracy: 1.0000

```

```

model.evaluate(X_test_resized, y_test)

```

```

8/8 [=====] - 3s 215ms/step - loss: 8.7594 - accuracy: 0.0000e+00
[8.759380340576172, 0.0]

```

2.4) EmoDB

```

Epoch 3/10
9/9 [=====] - 6s 663ms/step - loss: 1.1062 - accuracy: 0.6367
Epoch 4/10
9/9 [=====] - 6s 661ms/step - loss: 0.6534 - accuracy: 0.7678
Epoch 5/10
9/9 [=====] - 6s 662ms/step - loss: 0.3835 - accuracy: 0.8914
Epoch 6/10
9/9 [=====] - 6s 662ms/step - loss: 0.3716 - accuracy: 0.8689
Epoch 7/10
9/9 [=====] - 6s 662ms/step - loss: 0.2297 - accuracy: 0.9213
Epoch 8/10
9/9 [=====] - 6s 661ms/step - loss: 0.1096 - accuracy: 0.9775
Epoch 9/10
9/9 [=====] - 6s 664ms/step - loss: 0.1170 - accuracy: 0.9850
Epoch 10/10
9/9 [=====] - 6s 659ms/step - loss: 0.2414 - accuracy: 0.9251

model.evaluate(X_test_resized, y_test)

9/9 [=====] - 4s 304ms/step - loss: 7.2902 - accuracy: 0.0000e+00
[7.290168285369873, 0.0]

```

Looking at the complexity of the model and the limitations of google colab, I have reduced the input size for the model,i.e., I have taken 2000 training data points and 2000 testing data points.

3) Recurrent Neural Networks (RNN)

3.1) CIFAR-10

```
Epoch 3/10
200/200 [=====] - 111s 557ms/step - loss: 2.0085 - accuracy: 0.2645
Epoch 4/10
200/200 [=====] - 112s 558ms/step - loss: 1.9649 - accuracy: 0.2771
Epoch 5/10
200/200 [=====] - 111s 557ms/step - loss: 1.9583 - accuracy: 0.2816
Epoch 6/10
200/200 [=====] - 111s 557ms/step - loss: 1.9388 - accuracy: 0.2896
Epoch 7/10
200/200 [=====] - 111s 557ms/step - loss: 1.9371 - accuracy: 0.2899
Epoch 8/10
200/200 [=====] - 111s 556ms/step - loss: 1.9254 - accuracy: 0.2989
Epoch 9/10
200/200 [=====] - 111s 557ms/step - loss: 1.9188 - accuracy: 0.2966
Epoch 10/10
200/200 [=====] - 111s 556ms/step - loss: 1.9341 - accuracy: 0.2930
```

```
model.evaluate(test_images, test_labels)
```

```
157/157 [=====] - 38s 225ms/step - loss: 1.9601 - accuracy: 0.2912
[1.9600898027420044, 0.29120001196861267]
```

3.2) MNIST

```
print('Test Accuracy of the model on the 10000 test images: {} %'.format(100 * correct / total))

Test Accuracy of the model on the 10000 test images: 97.77 %
```

3.3) SAVEE

```
[ ] model.evaluate(X_test, y_test)
```

```
6/6 [=====] - 2s 57ms/step - loss: 1.4087 - accuracy: 0.4323
[1.408677577972412, 0.4322916567325592]
```

3.4) EmoDB

```
8/8 [=====] - 1s 130ms/step - loss: 0.2239 - accuracy: 0.9144
Epoch 50/50
8/8 [=====] - 1s 127ms/step - loss: 0.2858 - accuracy: 0.8984
```

```
[ ] model.evaluate(X_test, y_test)
```

```
6/6 [=====] - 2s 54ms/step - loss: 1.7593 - accuracy: 0.5590
[1.7592660188674927, 0.5590062141418457]
```

4) AlexNet

4.1) CIFAR-10

```
Epoch 1/5
16/16 [=====] - 61s 4s/step - loss: 2.6788 - accuracy: 0.1240
Epoch 2/5
16/16 [=====] - 59s 4s/step - loss: 2.6332 - accuracy: 0.0960
Epoch 3/5
16/16 [=====] - 60s 4s/step - loss: 2.5360 - accuracy: 0.1220
Epoch 4/5
16/16 [=====] - 59s 4s/step - loss: 2.4222 - accuracy: 0.1400
Epoch 5/5
16/16 [=====] - 60s 4s/step - loss: 2.3684 - accuracy: 0.1400
```

[+ Code](#)[+ Text](#)

```
[ ] model.evaluate(X_test_resized, y_test)
```

```
7/7 [=====] - 6s 753ms/step - loss: 2.3611 - accuracy: 0.0750
[2.3610661029815674, 0.07500000298023224]
```

4.2) MNIST

```
Epoch 1/5
63/63 [=====] - 474s 8s/step - loss: 2.0734 - accuracy: 0.2610
Epoch 2/5
63/63 [=====] - 476s 8s/step - loss: 1.7821 - accuracy: 0.3680
Epoch 3/5
63/63 [=====] - 468s 7s/step - loss: 1.6773 - accuracy: 0.4395
Epoch 4/5
63/63 [=====] - 469s 7s/step - loss: 1.5820 - accuracy: 0.4810
Epoch 5/5
63/63 [=====] - 472s 7s/step - loss: 1.5318 - accuracy: 0.5040
```

```
[ ] model.evaluate(X_test_new, y_test)
```

```
63/63 [=====] - 107s 2s/step - loss: 2.3417 - accuracy: 0.1170
[2.3417277336120605, 0.11699999868869781]
```

4.3) SAVEE


```
Epoch 4/10
8/8 [=====] - 56s 7s/step - loss: 2.4215 - accuracy: 0.1583
Epoch 5/10
8/8 [=====] - 56s 7s/step - loss: 2.2042 - accuracy: 0.2333
Epoch 6/10
8/8 [=====] - 57s 7s/step - loss: 2.2080 - accuracy: 0.2042
Epoch 7/10
8/8 [=====] - 56s 7s/step - loss: 2.1114 - accuracy: 0.2792
Epoch 8/10
8/8 [=====] - 57s 7s/step - loss: 2.1120 - accuracy: 0.2542
Epoch 9/10
8/8 [=====] - 56s 7s/step - loss: 2.0292 - accuracy: 0.2583
Epoch 10/10
8/8 [=====] - 57s 7s/step - loss: 2.1150 - accuracy: 0.2417

model.evaluate(X_test_resized, y_test)

8/8 [=====] - 13s 2s/step - loss: 2.2758 - accuracy: 0.2375
[2.275780200958252, 0.23749999701976776]
```

4.4) EmoDB

```
11/11 [=====] - 75s 7s/step - loss: 2.2610 - accuracy: 0.1838
Epoch 6/10
11/11 [=====] - 75s 7s/step - loss: 2.1741 - accuracy: 0.2025
Epoch 7/10
11/11 [=====] - 77s 7s/step - loss: 2.0738 - accuracy: 0.2336
Epoch 8/10
11/11 [=====] - 76s 7s/step - loss: 2.0492 - accuracy: 0.2679
Epoch 9/10
11/11 [=====] - 76s 7s/step - loss: 2.0359 - accuracy: 0.2679
Epoch 10/10
11/11 [=====] - 76s 7s/step - loss: 1.9726 - accuracy: 0.3115

model.evaluate(X_test_resized, y_test)

7/7 [=====] - 12s 2s/step - loss: 2.1748 - accuracy: 0.2336
[2.1748006343841553, 0.23364485800266266]
```

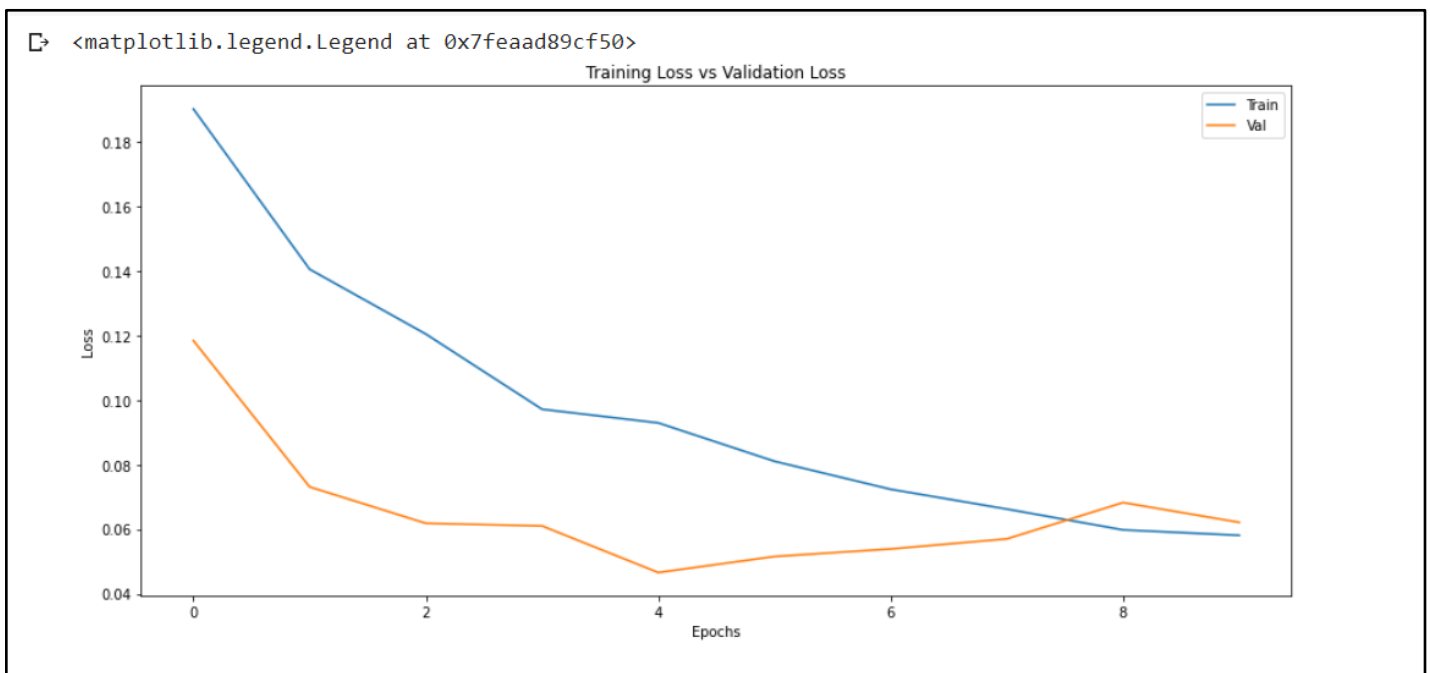
Looking at the complexity of the model and the limitations of google colab, I have reduced the input size for the model,i.e., I have taken 2000 training data points and 2000 testing data points.

5) GoogLeNet

5.1) CIFAR-10

```
output_1_loss: 2.1621 - val_output_accuracy: 0.2305 - val_auxilliary_output_1_accuracy: 0.2250 - val_auxilliary_output_2_accuracy: 0.2205  
  
output_2_loss: 2.0650 - val_output_accuracy: 0.2305 - val_auxilliary_output_1_accuracy: 0.2400 - val_auxilliary_output_2_accuracy: 0.2240  
  
output_2_loss: 2.0244 - val_output_accuracy: 0.2470 - val_auxilliary_output_1_accuracy: 0.2630 - val_auxilliary_output_2_accuracy: 0.2585  
  
output_2_loss: 2.0076 - val_output_accuracy: 0.2355 - val_auxilliary_output_1_accuracy: 0.2735 - val_auxilliary_output_2_accuracy: 0.2660
```

5.2) MNIST

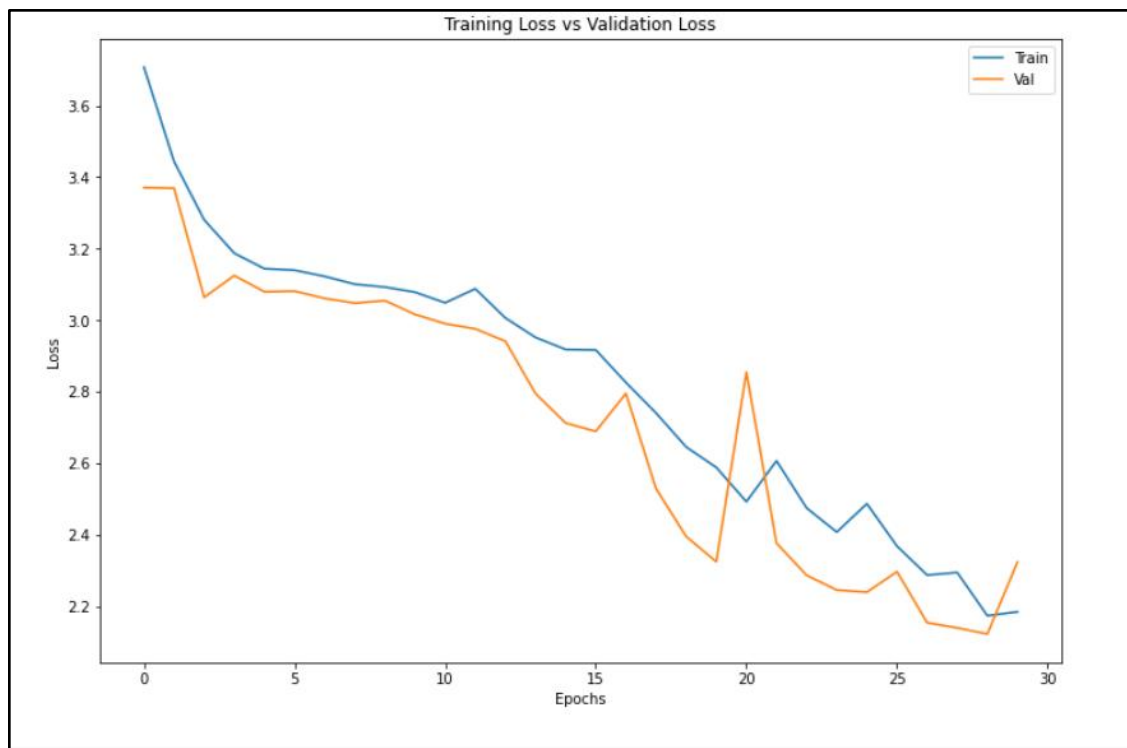




```
model.evaluate(x_test, y_test)
```

```
313/313 [=====] - 24s 6  
[0.05425573140382767,  
 0.03818700090050697,  
 0.02569129876792431,  
 0.027871087193489075,  
 0.9873999953269958,  
 0.9926000237464905,  
 0.9902999997138977]
```

5.3) SAVEE



```
0 5 10
model.evaluate(X_test, y_test)

6/6 [=====]
[2.324364423751831,
 1.458065390586853,
 1.4511628150939941,
 1.4365006685256958,
 0.3541666567325592,
 0.3333333432674408,
 0.3802083432674408]
```

5.4) EmoDB



```
model.evaluate(X_test, y_test)
```

```
7/7 [=====]
[2.8008506298065186,
 1.7843737602233887,
 1.6508854627609253,
 1.7373706102371216,
 0.30841121077537537,
 0.38785046339035034,
 0.3644859790802002]
```

Hidden Markov Model (HMM)

Dataset	Classifier	Train-Test Ratio	Precision	Recall	F1-Score	Support	Accuracy
Wine Dataset	GaussianHMM(With Tuning)	70-30	0.82	0.84	0.83	106	83
	GaussianHMM(With Tuning)	60-40	0.8	0.81	0.8	141	0.8
	GaussianHMM(With Tuning)	50-50	0.74	0.75	0.74	176	0.75
	GaussianHMM(With Tuning)	40-60	0.68	0.69	0.68	211	0.69
	GaussianHMM(With Tuning)	30-70	0.49	0.5	0.49	246	57
	GaussianHMM(Without Tuning)	70-30	0.91	0.95	0.93	54	92
	GaussianHMM(Without Tuning)	60-40	0.97	0.98	0.97	72	97

	GaussianHMM(Without Tuning)	50-50	0.34	0.35	0.34	89	35
	GaussianHMM(Without Tuning)	40-60	0.95	0.95	0.95	107	94
	GaussianHMM(Without Tuning)	30-70	0.33	0.39	0.36	125	37
	GMMHMM(With Tuning)	70-30	0.82	0.84	0.83	106	83
	GMMHMM(With Tuning)	60-40	0.8	0.81	0.8	141	0.8
	GMMHMM(With Tuning)	50-50	0.74	0.75	0.74	176	0.75
	GMMHMM(With Tuning)	40-60	0.68	0.69	0.68	211	0.69
	GMMHMM(With Tuning)	30-70	0.49	0.5	0.49	246	57
	GMMHMM(Without Tuning)	70-30	0.91	0.95	0.93	54	92
	GMMHMM(Without Tuning)	60-40	0.97	0.98	0.97	72	97
	GMMHMM(Without Tuning)	50-50	0.95	0.95	0.95	89	94
	GMMHMM(Without Tuning)	40-60	0.94	0.94	0.94	107	93
	GMMHMM(Without Tuning)	30-70	0.05	0.04	0.05	125	4.8
	MultinomialHMM(With Tuning)	70-30	0.82	0.84	0.83	106	83
	MultinomialHMM(With Tuning)	60-40	0.83	0.84	0.83	141	83
	MultinomialHMM(With Tuning)	50-50	0.74	0.75	0.74	176	75
	MultinomialHMM(With Tuning)	40-60	0.69	0.69	0.69	211	71

	MultinomialHMM(With Tuning)	30-70	0.52	0.51	0.43	246	43
	MultinomialHMM(Without Tuning)	70-30	0.82	0.84	0.83	106	83
	MultinomialHMM(Without Tuning)	60-40	0.83	0.84	0.83	141	83
	MultinomialHMM(Without Tuning)	50-50	0.74	0.75	0.74	176	75
	MultinomialHMM(Without Tuning)	40-60	0.69	0.69	0.69	211	71
	MultinomialHMM(Without Tuning)	30-70	0.52	0.51	0.43	246	0.43
Ionosphere Dataset	GaussianHMM(With Tuning)	70-30	0.82	0.84	0.83	106	83
	GaussianHMM(With Tuning)	60-40	0.8	0.81	0.8	141	0.8
	GaussianHMM(With Tuning)	50-50	0.74	0.75	0.74	176	0.75
	GaussianHMM(With Tuning)	40-60	0.68	0.69	0.68	211	0.69
	GaussianHMM(With Tuning)	30-70	0.49	0.5	0.49	246	57
	GaussianHMM(Without Tuning)	70-30	0.82	0.84	0.83	106	83
	GaussianHMM(Without Tuning)	60-40	0.81	0.82	0.81	141	81
	GaussianHMM(Without Tuning)	50-50	0.74	0.75	0.74	176	75
	GaussianHMM(Without Tuning)	40-60	0.68	0.69	0.68	211	69
	GaussianHMM(Without Tuning)	30-70	0.75	0.78	0.73	246	73
	GMMHMM(With Tuning)	70-30	0.82	0.84	0.83	106	83

	GMMHMM(With Tuning)	60-40	0.83	0.84	0.83	141	83
	GMMHMM(With Tuning)	50-50	0.74	0.75	0.74	176	75
	GMMHMM(With Tuning)	40-60	0.69	0.69	0.69	211	71
	GMMHMM(With Tuning)	30-70	0.52	0.51	0.43	246	43
	GMMHMM(Without Tuning)	70-30	0.82	0.84	0.83	106	83
	GMMHMM(Without Tuning)	60-40	0.83	0.84	0.83	141	83
	GMMHMM(Without Tuning)	50-50	0.74	0.75	0.74	176	75
	GMMHMM(Without Tuning)	40-60	0.69	0.69	0.69	211	71
	GMMHMM(Without Tuning)	30-70	0.52	0.51	0.43	246	0.43
	MultinomialHMM(With Tuning)	70-30	0.82	0.84	0.83	106	83
	MultinomialHMM(With Tuning)	60-40	0.83	0.84	0.83	141	83
	MultinomialHMM(With Tuning)	50-50	0.74	0.75	0.74	176	75
	MultinomialHMM(With Tuning)	40-60	0.69	0.69	0.69	211	71
	MultinomialHMM(With Tuning)	30-70	0.52	0.51	0.43	246	43
	MultinomialHMM(Without Tuning)	70-30	0.82	0.84	0.83	106	83
	MultinomialHMM(Without Tuning)	60-40	0.83	0.84	0.83	141	83
	MultinomialHMM(Without Tuning)	50-50	0.74	0.75	0.74	176	75

	MultinomialHMM(Without Tuning)	40-60	0.69	0.69	0.69	211	71
	MultinomialHMM(Without Tuning)	30-70	0.52	0.51	0.43	246	0.43
Breast Cancer Dataset	GaussianHMM(With Tuning)	70-30	0.94	0.95	0.94	171	94
	GaussianHMM(With Tuning)	60-40	0.94	0.95	0.94	228	94
	GaussianHMM(With Tuning)	50-50	0.07	0.06	0.06	285	6
	GaussianHMM(With Tuning)	40-60	0.85	0.84	0.84	342	86
	GaussianHMM(With Tuning)	30-70	0.91	0.91	0.91	399	91
	GaussianHMM(Without Tuning)	70-30	0.94	0.96	0.95	171	95
	GaussianHMM(Without Tuning)	60-40	0.92	0.93	0.92	228	92
	GaussianHMM(Without Tuning)	50-50	0.93	0.94	0.93	285	93
	GaussianHMM(Without Tuning)	40-60	0.85	0.84	0.84	342	86
	GaussianHMM(Without Tuning)	30-70	0.91	0.91	0.91	399	91
	GMMHMM(With Tuning)	70-30	0.92	0.93	0.92	171	92
	GMMHMM(With Tuning)	60-40	0.91	0.91	0.91	228	91
	GMMHMM(With Tuning)	50-50	0.91	0.92	0.91	285	91
	GMMHMM(With Tuning)	40-60	0.89	0.91	0.9	342	90
	GMMHMM(With Tuning)	30-70	0.9	0.78	0.8	399	0.83

	GMMHMM(Without Tuning)	70-30	0.92	0.93	0.92	171	92
	GMMHMM(Without Tuning)	60-40	0.91	0.91	0.91	228	91
	GMMHMM(Without Tuning)	50-50	0.91	0.92	0.91	285	91
	GMMHMM(Without Tuning)	40-60	0.89	0.91	0.9	342	90
	GMMHMM(Without Tuning)	30-70	0.9	0.91	0.9	399	90
	MultinomialHMM(With Tuning)	70-30	0.51	0.51	0.51	171	57
	MultinomialHMM(With Tuning)	60-40	0.54	0.54	0.54	228	59
	MultinomialHMM(With Tuning)	50-50	0.54	0.54	0.54	285	57
	MultinomialHMM(With Tuning)	40-60	0.53	0.53	0.53	342	58
	MultinomialHMM(With Tuning)	30-70	0.54	0.54	0.54	399	57
	MultinomialHMM(Without Tuning)	70-30	0.51	0.51	0.51	171	57
	MultinomialHMM(Without Tuning)	60-40	0.54	0.54	0.54	228	59
	MultinomialHMM(Without Tuning)	50-50	0.54	0.54	0.54	285	57
	MultinomialHMM(Without Tuning)	40-60	0.53	0.53	0.53	342	58
	MultinomialHMM(Without Tuning)	30-70	0.54	0.54	0.54	399	57

Convolutional Neural Networks(CNN)							
Dataset	Accuracy						
CIFAR-10	68						
MNIST	99						
SAVEE	29						
EmoDB	50						
Other Deep Learning Models							
Models	Dataset	Accura cy					
VGG-16	CIFAR-10	9.8					
	MNIST	10.95					
	SAVEE	12.92					
	EmoDB	25					
ResNet-50	CIFAR-10	27					
	MNIST	99					
	SAVEE	99					
	EmoDB	92					
Recurrent Neural Networks (RNN)	CIFAR-10	29					
	MNIST	97					
	SAVEE	43					
	EmoDB	55					
AlexNet	CIFAR-10	7.5					
	MNIST	11.69					

	SAVEE	23.74					
	EmoDB	23.36					
GoogLeNet	CIFAR-10	26.6					
	MNIST	99					
	SAVEE	38					
	EmoDB	36					

WINE DATASET

GaussianHMM(Without Tuning)[70-30 split]

import pandas as pd

import numpy as np

Dataset Preparation

df = pd.read_csv('wine.data',header=None)

**col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of
ash','Magnesium','Total phenols','Flavanoids',
'Nonflavanoid phenols','Proanthocyanins','Color
intensity','Hue','OD280/OD315 of diluted wines','Proline']**

df.columns = col_name

X = df.drop(['Class'], axis=1)

y = df['Class']

from sklearn.model_selection import train_test_split

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=3, covariance_type="full")
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 0:
```

```
        strings[i] = 1
```

```
    elif y_pred[i] == 1:
```

```
        strings[i] = 2
```

```
    else:
```

```
        strings[i] = 3
```

```
strings = strings.astype(np.int)
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')  
plt.show()
```

```
# WINE DATASET  
# GaussianHMM(Without Tuning)[60-40 split]
```

```
import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv('wine.data',header=None)

col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of
ash','Magnesium','Total phenols','Flavanoids',
            'Nonflavanoid phenols','Proanthocyanins','Color
intensity','Hue','OD280/OD315 of diluted wines','Proline']

df.columns = col_name

X = df.drop(['Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=3, covariance_type="full")  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 0:  
        strings[i] = 1  
    elif y_pred[i] == 1:  
        strings[i] = 2  
    else:  
        strings[i] = 3
```

```
strings = strings.astype(np.int)
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```



```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# WINE DATASET  
# GaussianHMM(Without Tuning)[50-50 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv("wine.data", header=None)
```

```
col_name = ['Class', 'Alcohol', 'Malic acid', 'Ash', 'Alcalinity of  
ash', 'Magnesium', 'Total phenols', 'Flavanoids',  
            'Nonflavanoid phenols', 'Proanthocyanins', 'Color  
intensity', 'Hue', 'OD280/OD315 of diluted wines', 'Proline']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=None)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=3, covariance_type="full")  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 0:
```

```
strings[i] = 1
elif y_pred[i] == 1:
strings[i] = 2
else:
strings[i] = 3
```

```
strings = strings.astype(np.int)
```

```
from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score
```

```
print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
```

```
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# WINE DATASET
```

```
# GaussianHMM(Without Tuning)[40-60 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv("wine.data", header=None)
```

```
col_name = ['Class', 'Alcohol', 'Malic acid', 'Ash', 'Alcalinity of
ash', 'Magnesium', 'Total phenols', 'Flavanoids',
            'Nonflavanoid phenols', 'Proanthocyanins', 'Color
intensity', 'Hue', 'OD280/OD315 of diluted wines', 'Proline']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
train_test_split(X, y, train_size=0.4, test_size=0.6, random_state=10)
```

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

X_test = sc.transform(X_test)

Classification

from hmmlearn import hmm

classifier = hmm.GaussianHMM(n_components=3, covariance_type="full")

classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)

strings = np.empty(size, np.unicode_)

for i in range (size):

if y_pred[i] == 0:

strings[i] = 1

elif y_pred[i] == 1:

strings[i] = 2

else:

strings[i] = 3

strings = strings.astype(np.int)

from sklearn.metrics import classification_report, confusion_matrix,

accuracy_score

```
print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# WINE DATASET
# GaussianHMM(Without Tuning)[30-70 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('wine.data',header=None)
```

```
col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of  
ash','Magnesium','Total phenols','Flavanoids',  
            'Nonflavanoid phenols','Proanthocyanins','Color  
intensity','Hue','OD280/OD315 of diluted wines','Proline']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=3, covariance_type="full")  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 0:
```

```
        strings[i] = 1
```

```
    elif y_pred[i] == 1:
```

```
        strings[i] = 2
```

```
    else:
```

```
        strings[i] = 3
```

```
strings = strings.astype(np.int)
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```



```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# WINE DATASET
# GaussianHMM(With Tuning)[70-30 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('wine.data', header=None)
```

```
col_name = ['Class', 'Alcohol', 'Malic acid', 'Ash', 'Alcalinity of
ash', 'Magnesium', 'Total phenols', 'Flavanoids',
            'Nonflavanoid phenols', 'Proanthocyanins', 'Color
intensity', 'Hue', 'OD280/OD315 of diluted wines', 'Proline']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=3, covariance_type="full")  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 0:  
        strings[i] = 1  
    elif y_pred[i] == 1:  
        strings[i] = 2  
    else:
```

```
strings[i] = 3
```

```
strings = strings.astype(np.int)
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
#####  
#####  
#####  
#####  
#####  
#####
```

WINE DATASET

GMMHMM(Without Tuning)[70-30 split]

import pandas as pd

import numpy as np

Dataset Preparation

df = pd.read_csv('wine.data',header=None)

**col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of
ash','Magnesium','Total phenols','Flavanoids',
 'Nonflavanoid phenols','Proanthocyanins','Color
intensity','Hue','OD280/OD315 of diluted wines','Proline']**

df.columns = col_name

X = df.drop(['Class'], axis=1)

y = df['Class']

from sklearn.model_selection import train_test_split

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=3,
```

```
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 0:
```

```
        strings[i] = 3
```

```
    elif y_pred[i] == 1:
```

```
        strings[i] = 2
```

```
    else:
```

```
        strings[i] = 1
```

```
strings = strings.astype(np.int)
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# WINE DATASET  
# GMMHMM(Without Tuning)[60-40 split]
```

```
import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv('wine.data',header=None)

col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of
ash','Magnesium','Total phenols','Flavanoids',
            'Nonflavanoid phenols','Proanthocyanins','Color
intensity','Hue','OD280/OD315 of diluted wines','Proline']

df.columns = col_name

X = df.drop(['Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)
```

```
# Classification
# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.GMMHMM(n_components=3,
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)
strings = np.empty(size, np.unicode_)

for i in range (size):
    if y_pred[i] == 0:
        strings[i] = 3
    elif y_pred[i] == 1:
        strings[i] = 1
    else:
        strings[i] = 2

strings = strings.astype(np.int)

from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))

print("-----")
print("-----")
```



```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# WINE DATASET  
# GMMHMM(Without Tuning)[50-50 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv("wine.data", header=None)
```

```
col_name = ['Class', 'Alcohol', 'Malic acid', 'Ash', 'Alcalinity of  
ash', 'Magnesium', 'Total phenols', 'Flavanoids',
```

**'Nonflavanoid phenols','Proanthocyanins','Color
intensity','Hue','OD280/OD315 of diluted wines','Proline']**

df.columns = col_name

**X = df.drop(['Class'], axis=1)
y = df['Class']**

from sklearn.model_selection import train_test_split

**X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=10)**

Feature Scaling

from sklearn.preprocessing import StandardScaler

**sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)**

Classification

**# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.GMMHMM(n_components=3,
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
classifier.fit(X_train)**

y_pred = classifier.predict(X_test)

```
size = len(y_pred)
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
    if y_pred[i] == 0:
        strings[i] = 1
    elif y_pred[i] == 1:
        strings[i] = 3
    else:
        strings[i] = 2
```

```
strings = strings.astype(np.int)
```

```
from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score
```

```
print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')  
plt.show()
```

```
# WINE DATASET  
# GMMHMM(Without Tuning)[40-60 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv('wine.data', header=None)
```

```
col_name = ['Class', 'Alcohol', 'Malic acid', 'Ash', 'Alcalinity of  
ash', 'Magnesium', 'Total phenols', 'Flavanoids',  
            'Nonflavanoid phenols', 'Proanthocyanins', 'Color  
intensity', 'Hue', 'OD280/OD315 of diluted wines', 'Proline']
```

```
df.columns = col_name
```

```
X = df.drop(['Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=3,  
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 0:
```

```
        strings[i] = 3
```

```
    elif y_pred[i] == 1:
```

```
        strings[i] = 2
```

```
    else:
```

```
        strings[i] = 1
```

```
strings = strings.astype(np.int)
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# WINE DATASET
# GMMHMM(Without Tuning)[30-70 split]

import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv('wine.data',header=None)

col_name = ['Class','Alcohol','Malic acid','Ash','Alcalinity of
ash','Magnesium','Total phenols','Flavanoids',
            'Nonflavanoid phenols','Proanthocyanins','Color
intensity','Hue','OD280/OD315 of diluted wines','Proline']

df.columns = col_name

X = df.drop(['Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.GMMHMM(n_components=3,
random_state=10,covariance_type='diag',algorithm='viterbi',n_iter=10)
classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)
strings = np.empty(size, np.unicode_)

for i in range (size):
    if y_pred[i] == 0:
        strings[i] = 3
    elif y_pred[i] == 1:
        strings[i] = 1
    else:
        strings[i] = 2

strings = strings.astype(np.int)

from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```



```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# IONOSPHERE DATASET  
# GaussianHMM(Without Tuning)[70-30 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv('ionosphere.data',header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19']
```

```
, '20', '21', '22', '23', '24', '25', '26', '27', '28', '29', '30', '31', '32', '33', '34', 'Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1', '2', 'Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
```

```
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2, covariance_type="full")
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ("g")  
    else:  
        strings[i] = ("b")
```

strings

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
plt.show()
```

```
# IONOSPHERE DATASET
# GaussianHMM(Without Tuning)[60-40 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('ionosphere.data', header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2, covariance_type="full")
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ("g")
```

```
    else:
```

```
        strings[i] = ("b")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,
```

```
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# IONOSPHERE DATASET  
# GaussianHMM(Without Tuning)[50-50 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv('ionosphere.data', header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19']
```

```
, '20', '21', '22', '23', '24', '25', '26', '27', '28', '29', '30', '31', '32', '33', '34', 'Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1', '2', 'Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
```

```
train_test_split(X, y, train_size=0.5, test_size=0.5, random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2, covariance_type="full")
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ("g")  
    else:  
        strings[i] = ("b")
```

strings

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```



```
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
plt.show()
```

```
# IONOSPHERE DATASET
# GaussianHMM(Without Tuning)[40-60 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('ionosphere.data', header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,
```

```
covariance_type='full',algorithm='viterbi')
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ("g")
```

```
    else:
```

```
        strings[i] = ("b")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,
```

```
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
cm = confusion_matrix(y_test, strings)
```

```
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
```

```
plt.show()
```

```
# IONOSPHERE DATASET
```

```
# GaussianHMM(Without Tuning)[30-70 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('ionosphere.data', header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,  
covariance_type='diag',algorithm='viterbi')  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ('g')
    else:
        strings[i] = ('b')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score
```

```
print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# IONOSPHERE DATASET
```

```
# GaussianHMM(With Tuning)[70-30 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('ionosphere.data',header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

X_test = sc.transform(X_test)

Classification

from hmmlearn import hmm

**classifier = hmm.GaussianHMM(n_components=2,
covariance_type='full',n_iter=5,algorithm='viterbi',verbose=False)
classifier.fit(X_train)**

y_pred = classifier.predict(X_test)

size = len(y_pred)

strings = np.empty(size, np.unicode_)

for i in range (size):

if y_pred[i] == 1:

strings[i] = ('g')

else:

strings[i] = ('b')

strings

**from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score**

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')  
plt.show()
```

```
# IONOSPHERE DATASET  
# GaussianHMM(With Tuning)[60-40 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation
```



```
df = pd.read_csv('ionosphere.data',header=None)

col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']

df.columns = col_name

X = df.drop(['1','2','Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
from hmmlearn import hmm

classifier = hmm.GaussianHMM(n_components=2,
covariance_type='full',n_iter=2,algorithm='viterbi',verbose=False)
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ("g")
```

```
    else:
```

```
        strings[i] = ("b")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# IONOSPHERE DATASET  
# GaussianHMM(With Tuning)[50-50 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv('ionosphere.data', header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
            , '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,  
covariance_type='full',n_iter=5,algorithm='viterbi',verbose=False)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ('b')
```

```
    else:
```

```
        strings[i] = ('g')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# IONOSPHERE DATASET  
# GaussianHMM(With Tuning)[40-60 split]
```

```
import pandas as pd
```

```
import numpy as np

# Dataset Preparation
df = pd.read_csv('ionosphere.data',header=None)

col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']

df.columns = col_name

X = df.drop(['1','2','Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,  
covariance_type='full',n_iter=5,algorithm='viterbi',verbose=False)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ('g')  
    else:  
        strings[i] = ('b')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# IONOSPHERE DATASET
# GaussianHMM(With Tuning)[30-70 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('ionosphere.data', header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
y = df['Class']
```



```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,  
covariance_type='full',n_iter=5,algorithm='viterbi',verbose=False)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ("b")
```

```
    else:
```

```
        strings[i] = ("g")
```

strings

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
#####  
#####
```

```
import pandas as pd
```

```
import numpy as np

# Dataset Preparation
df = pd.read_csv('ionosphere.data',header=None)

col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']

df.columns = col_name

X = df.drop(['1','2','Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
# from hmmlearn import hmm
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=2,  
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ('g')  
    else:  
        strings[i] = ('b')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# IONOSPHERE DATASET
# GMMHMM(Without Tuning)[60-40 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('ionosphere.data', header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=2,  
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ('g')
```

```
    else:
```

```
        strings[i] = ('b')
```

strings

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# IONOSPHERE DATASET  
# GMMHMM(Without Tuning)[50-50 split]
```



```
import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv('ionosphere.data',header=None)

col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']

df.columns = col_name

X = df.drop(['1','2','Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
```

```
# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.GMMHMM(n_components=2,
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ("g")
    else:
        strings[i] = ("b")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score
```

```
print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
cm = confusion_matrix(y_test, strings)
```

```
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
```

```
plt.show()
```

```
# IONOSPHERE DATASET
```

```
# GMMHMM(Without Tuning)[40-60 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('ionosphere.data', header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.GMMHMM(n_components=2,
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ("g")
    else:
```

```
strings[i] = ('b')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

IONOSPHERE DATASET

GMMHMM(Without Tuning)[30-70 split]

import pandas as pd

import numpy as np

Dataset Preparation

df = pd.read_csv('ionosphere.data',header=None)

**col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
'20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']**

df.columns = col_name

X = df.drop(['1','2','Class'], axis=1)

y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =

train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

X_test = sc.transform(X_test)

```
# Classification
# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.GMMHMM(n_components=2,
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)
strings = np.empty(size, np.unicode_)

for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ('g')
    else:
        strings[i] = ('b')

strings

from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))

print("-----")
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# IONOSPHERE DATASET  
# GMMHMM(With Tuning)[70-30 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv("ionosphere.data", header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
            , '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```



```
X = df.drop(['1','2','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm  
import hmmlearn  
classifier = hmmlearn.hmm.GMMHMM(n_components=2,  
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:
```

```
strings[i] = ("g")
else:
strings[i] = ("b")
```

strings

```
from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score
```

```
print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
```

```
plt.show()
```

```
# IONOSPHERE DATASET
```

```
# GMMHMM(With Tuning)[60-40 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('ionosphere.data',header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
```

```
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```

X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.GMMHMM(n_components=2,
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)
strings = np.empty(size, np.unicode_)

for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ("g")
    else:
        strings[i] = ("b")

strings

from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))

print("-----")

```

```
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# IONOSPHERE DATASET  
# GMMHMM(With Tuning)[50-50 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv('ionosphere.data', header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
            , '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=2,  
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ('g')  
    else:  
        strings[i] = ('b')
```

strings

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns
```

```
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# IONOSPHERE DATASET
```

```
# GMMHMM(With Tuning)[40-60 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('ionosphere.data',header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
```

```
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```



```
sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.GMMHMM(n_components=2,
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)
strings = np.empty(size, np.unicode_)

for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ("g")
    else:
        strings[i] = ("b")

strings

from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# IONOSPHERE DATASET
# GMMHMM(With Tuning)[30-70 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('ionosphere.data', header=None)
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19']
```

```
,'20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
```

```
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=2,
```

```
random_state=10,covariance_type='full',algorithm='viterbi',n_iter=10)
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ("g")  
    else:  
        strings[i] = ("b")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

[illegible]

MultinomialHMM(Without Tuning)[70-30 split]

```
import pandas as pd
import numpy as np

# Dataset Preparation
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']

df.columns = col_name

X = df.drop(['1','2','Class'], axis=1)
y = df['Class']

X = df.drop(['1','Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)
```

```
# Classification
# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,
random_state=15,n_iter=10,algorithm='viterbi',params='ste')
```

```
import math
row = len(X_train)
col = len(X_train[0])
new = [1] * 33
for i in range(row):
    for j in range(col):
        X_train[i][j] = X_train[i][j]*10
        X_train[i][j] = math.floor(X_train[i][j])
        x = X_train[i].astype(np.int)
        new = np.vstack([new,x])
```

```
y = new
y = np.absolute(y)
X_train = y
```

```
import math
row = len(X_test)
col = len(X_test[0])
new
for i in range(row):
    for j in range(col):
        X_test[i][j] = X_test[i][j]*10
        X_test[i][j] = math.floor(X_test[i][j])
        x = X_test[i].astype(np.int)
        new = np.vstack([new,x])
```

```
y = new
```

```
y = np.absolute(y)
```

```
X_test = y
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ('g')
```

```
    else:
```

```
        strings[i] = ('b')
```

```
strings
```

```
strings = strings[0:106]
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```



```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# IONOSPHERE DATASET  
# MultinomialHMM(Without Tuning)[60-40 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
            , '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)  
y = df['Class']
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,  
random_state=15,n_iter=10,algorithm='viterbi',params='ste')
```

```
import math
```

```
row = len(X_train)
```

```
col = len(X_train[0])
```

```
new = [1] * 33
```

```
for i in range(row):
```

```
    for j in range(col):
```

```
X_train[i][j] = X_train[i][j]*10  
X_train[i][j] = math.floor(X_train[i][j])  
x = X_train[i].astype(np.int)  
new = np.vstack([new,x])
```

```
y = new  
y = np.absolute(y)  
X_train = y
```

```
import math  
row = len(X_test)  
col = len(X_test[0])  
new  
for i in range(row):  
    for j in range(col):  
        X_test[i][j] = X_test[i][j]*10  
        X_test[i][j] = math.floor(X_test[i][j])  
        x = X_test[i].astype(np.int)  
        new = np.vstack([new,x])
```

```
y = new  
y = np.absolute(y)  
X_test = y
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ("g")  
    else:  
        strings[i] = ("b")
```

```
strings  
strings = strings[0:141]
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
plt.show()
```

```
# IONOSPHERE DATASET
```

```
# MultinomialHMM(Without Tuning)[50-50 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
```

```
y = df['Class']
```

```
X = df.drop(['1','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
```

```
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=10)
```

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

X_test = sc.transform(X_test)

Classification

from hmmlearn import hmm

import hmmlearn

**classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,
random_state=15,n_iter=10,algorithm='viterbi',params='ste')**

import math

row = len(X_train)

col = len(X_train[0])

new = [1] * 33

for i in range(row):

for j in range(col):

X_train[i][j] = X_train[i][j]*10

X_train[i][j] = math.floor(X_train[i][j])

x = X_train[i].astype(np.int)

new = np.vstack([new,x])

y = new

y = np.absolute(y)

X_train = y

import math

row = len(X_test)

```
col = len(X_test[0])
new
for i in range(row):
    for j in range(col):
        X_test[i][j] = X_test[i][j]*10
        X_test[i][j] = math.floor(X_test[i][j])
    x = X_test[i].astype(np.int)
    new = np.vstack([new,x])
```

```
y = new
y = np.absolute(y)
X_test = y
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ("b")
    else:
        strings[i] = ("g")
```

```
strings
strings = strings[0:176]
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')  
plt.show()
```

```
# IONOSPHERE DATASET  
# MultinomialHMM(Without Tuning)[40-60 split]
```

```
import pandas as pd  
import numpy as np
```


Dataset Preparation

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)  
y = df['Class']
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

Feature Scaling

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

Classification

```
# from hmmlearn import hmm  
import hmmlearn
```

```
classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,  
random_state=15,n_iter=10,algorithm='viterbi',params='ste')
```

```
import math  
row = len(X_train)  
col = len(X_train[0])  
new = [1] * 33  
for i in range(row):  
    for j in range(col):  
        X_train[i][j] = X_train[i][j]*10  
        X_train[i][j] = math.floor(X_train[i][j])  
        x = X_train[i].astype(np.int)  
        new = np.vstack([new,x])
```

```
y = new  
y = np.absolute(y)  
X_train = y
```

```
import math  
row = len(X_test)  
col = len(X_test[0])  
new  
for i in range(row):  
    for j in range(col):  
        X_test[i][j] = X_test[i][j]*10  
        X_test[i][j] = math.floor(X_test[i][j])  
        x = X_test[i].astype(np.int)  
        new = np.vstack([new,x])
```

```
y = new  
y = np.absolute(y)
```

```
X_test = y
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ('b')
```

```
    else:
```

```
        strings[i] = ('g')
```

```
strings
```

```
strings = strings[0:211]
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# IONOSPHERE DATASET
```

```
# MultinomialHMM(Without Tuning)[30-70 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
```

```
col_name = ['1','2','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
            '20','21','22','23','24','25','26','27','28','29','30','31','32','33','34','Class']
```

```
df.columns = col_name
```

```
X = df.drop(['1','2','Class'], axis=1)
y = df['Class']
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,  
random_state=15,n_iter=10,algorithm='viterbi',params='ste')
```

```
import math
```

```
row = len(X_train)
```

```
col = len(X_train[0])
```

```
new = [1] * 33
```

```
for i in range(row):
```

```
    for j in range(col):
```

```
        X_train[i][j] = X_train[i][j]*10
```

```
        X_train[i][j] = math.floor(X_train[i][j])
```

```
        x = X_train[i].astype(np.int)
```

```
new = np.vstack([new,x])
```

```
y = new
```

```
y = np.absolute(y)
```

```
X_train = y
```

```
import math
```

```
row = len(X_test)
```

```
col = len(X_test[0])
```

```
new
```

```
for i in range(row):
```

```
    for j in range(col):
```

```
        X_test[i][j] = X_test[i][j]*10
```

```
        X_test[i][j] = math.floor(X_test[i][j])
```

```
        x = X_test[i].astype(np.int)
```

```
        new = np.vstack([new,x])
```

```
y = new
```

```
y = np.absolute(y)
```

```
X_test = y
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
strings[i] = ('b')
else:
strings[i] = ('g')
```

```
strings
strings = strings[0:246]
```

```
from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score
```

```
print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
```

```
plt.show()
```

```
# BREAST CANCER DATASET
```

```
# GaussianHMM(Without Tuning)[70-30 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('wdbc.data',header=None)
```

```
col_name =
```

```
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
```

```
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```



```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2, covariance_type="full")
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ("M")
```

```
    else:
```

```
        strings[i] = ("B")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# BREAST CANCER DATASET  
# GaussianHMM(Without Tuning)[60-40 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =  
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2, covariance_type="full")  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ("M")  
    else:  
        strings[i] = ("B")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# BREAST CANCER DATASET
# GaussianHMM(Without Tuning)[50-50 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'
    ,'20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=10)
```

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

X_test = sc.transform(X_test)

Classification

from hmmlearn import hmm

classifier = hmm.GaussianHMM(n_components=2,

covariance_type="full",n_iter=10)

classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)

strings = np.empty(size, np.unicode_)

for i in range (size):

if y_pred[i] == 1:

strings[i] = ("M")

else:

strings[i] = ("B")

strings

from sklearn.metrics import classification_report, confusion_matrix,

accuracy_score

```

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))

print("-----")
print("-----")

print("Performance Evaluation")
print(classification_report(y_test, strings))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, strings))

import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()

# BREAST CANCER DATASET
# GaussianHMM(Without Tuning)[40-60 split]

import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv('wdbc.data', header=None)

```

```
col_name =  
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,  
covariance_type="full",n_iter=10)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```



```
size = len(y_pred)
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ('M')
    else:
        strings[i] = ('B')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score
```

```
print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# BREAST CANCER DATASET  
# GaussianHMM(Without Tuning)[30-70 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =  
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)
```

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

X_test = sc.transform(X_test)

Classification

from hmmlearn import hmm

classifier = hmm.GaussianHMM(n_components=2,

covariance_type="full",n_iter=10)

classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)

strings = np.empty(size, np.unicode_)

for i in range (size):

if y_pred[i] == 1:

strings[i] = ("M")

else:

strings[i] = ("B")

strings

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')  
plt.show()
```

```
# BREAST CANCER DATASET  
# GaussianHMM(With Tuning)[70-30 split]
```

```
import pandas as pd  
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv('wdbc.data',header=None)
```

```
col_name =
```

```
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
```

```
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

Feature Scaling

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

Classification

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,
```

```
covariance_type='full',n_iter=10,algorithm='viterbi',verbose=False)
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ('M')
```

```
    else:
```

```
        strings[i] = ('B')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
cm = confusion_matrix(y_test, strings)
```

```
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
```

```
plt.show()
```

```
# BREAST CANCER DATASET
```

```
# GaussianHMM(With Tuning)[60-40 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =
```

```
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
 '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,
```

```
covariance_type='full',n_iter=10,algorithm='viterbi',verbose=False)
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ("M")
```

```
    else:
```

```
        strings[i] = ("B")
```

```
strings
```



```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')  
plt.show()
```

```
# BREAST CANCER DATASET  
# GaussianHMM(With Tuning)[50-50 split]
```

```
import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv('wdbc.data',header=None)

col_name =
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'
    ,'20','21','22','23','24','25','26','27','28','29','30','31','32']

df.columns = col_name

X = df.drop(['1','Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,  
covariance_type='full',n_iter=10,algorithm='viterbi',verbose=False)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ('M')  
    else:  
        strings[i] = ('B')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# BREAST CANCER DATASET
# GaussianHMM(With Tuning)[40-60 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'
, '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,  
covariance_type='full',n_iter=10,algorithm='viterbi',verbose=False)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ('M')  
    else:  
        strings[i] = ('B')
```

strings

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# BREAST CANCER DATASET  
# GaussianHMM(With Tuning)[30-70 split]
```

```
import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv('wdbc.data',header=None)

col_name =
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'
    ,'20','21','22','23','24','25','26','27','28','29','30','31','32']

df.columns = col_name

X = df.drop(['1','Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
```

```
from hmmlearn import hmm
```

```
classifier = hmm.GaussianHMM(n_components=2,  
covariance_type='full',n_iter=10,algorithm='viterbi',verbose=False)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ('M')  
    else:  
        strings[i] = ('B')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")
```



```
print(classification_report(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print('Accuracy:')
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

cm = confusion_matrix(y_test, strings)

```
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
```

plt.show()

[illegible]


```
col_name =  
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=2, random_state=10)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ("M")  
    else:  
        strings[i] = ("B")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# BREAST CANCER DATASET
# GMMHMM(Without Tuning)[60-40 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'
    ,'20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)
```

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

X_test = sc.transform(X_test)

Classification

from hmmlearn import hmm

import hmmlearn

classifier = hmmlearn.hmm.GMMHMM(n_components=2, random_state=2)

classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)

strings = np.empty(size, np.unicode_)

for i in range (size):

if y_pred[i] == 1:

strings[i] = ("M")

else:

strings[i] = ("B")

strings

from sklearn.metrics import classification_report, confusion_matrix,

accuracy_score

print("Confusion Matrix:")

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
cm = confusion_matrix(y_test, strings)
```

```
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
```

```
plt.show()
```

```
# BREAST CANCER DATASET
```

```
# GMMHMM(Without Tuning)[50-50 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =  
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=2, random_state=10)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```



```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ("M")  
    else:  
        strings[i] = ("B")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# BREAST CANCER DATASET
# GMMHMM(Without Tuning)[40-60 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'
    ,'20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

X_test = sc.transform(X_test)

Classification

from hmmlearn import hmm

import hmmlearn

classifier = hmmlearn.hmm.GMMHMM(n_components=2, random_state=10)

classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)

strings = np.empty(size, np.unicode_)

for i in range (size):

if y_pred[i] == 1:

strings[i] = ("M")

else:

strings[i] = ("B")

strings

from sklearn.metrics import classification_report, confusion_matrix,

accuracy_score

print("Confusion Matrix:")

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
cm = confusion_matrix(y_test, strings)
```

```
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
```

```
plt.show()
```

```
# BREAST CANCER DATASET
```

```
# GMMHMM(Without Tuning)[30-70 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('wdbc.data',header=None)
```

```
col_name =  
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=2, random_state=2)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ("M")  
    else:  
        strings[i] = ("B")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()
```

```
# BREAST CANCER DATASET
# GMMHMM(With Tuning)[70-30 split]
```

```
import pandas as pd
import numpy as np
```

```
# Dataset Preparation
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'
    ,'20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

X_test = sc.transform(X_test)

Classification

from hmmlearn import hmm

import hmmlearn

classifier = hmmlearn.hmm.GMMHMM(n_components=2,

random_state=10,covariance_type='diag',algorithm='viterbi',n_iter=10)

classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)

strings = np.empty(size, np.unicode_)

for i in range (size):

if y_pred[i] == 1:

strings[i] = ("M")

else:

strings[i] = ("B")

strings

from sklearn.metrics import classification_report, confusion_matrix,

accuracy_score


```

print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))

print("-----")
print("-----")

print("Performance Evaluation")
print(classification_report(y_test, strings))

print("-----")
print("-----")

print("Accuracy:")
print(accuracy_score(y_test, strings))

import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
plt.show()

# BREAST CANCER DATASET
# GMMHMM(With Tuning)[60-40 split]

import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv('wdbc.data', header=None)

```

```
col_name =  
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()  
X_train = sc.fit_transform(X_train)  
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=2,  
random_state=2,covariance_type='diag',algorithm='viterbi',n_iter=10)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ('M')
    else:
        strings[i] = ('B')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score
```

```
print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# BREAST CANCER DATASET  
# GMMHMM(With Tuning)[50-50 split]
```

```
import pandas as pd  
import numpy as np
```

```
# Dataset Preparation  
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =  
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)  
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=10)
```

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

X_test = sc.transform(X_test)

Classification

from hmmlearn import hmm

import hmmlearn

**classifier = hmmlearn.hmm.GMMHMM(n_components=2,
random_state=10,covariance_type='diag',algorithm='viterbi',n_iter=10)**

classifier.fit(X_train)

y_pred = classifier.predict(X_test)

size = len(y_pred)

strings = np.empty(size, np.unicode_)

for i in range (size):

if y_pred[i] == 1:

strings[i] = ("M")

else:

strings[i] = ("B")

strings

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')  
plt.show()
```

```
# BREAST CANCER DATASET  
# GMMHMM(With Tuning)[40-60 split]
```

```
import pandas as pd  
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv('wdbc.data',header=None)
```

```
col_name =
```

```
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
```

```
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

Feature Scaling

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

Classification

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=2,
```

```
random_state=10,covariance_type='diag',algorithm='viterbi',n_iter=10)
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ('M')
```

```
    else:
```

```
        strings[i] = ('B')
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
```



```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
cm = confusion_matrix(y_test, strings)
```

```
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')
```

```
plt.show()
```

```
# BREAST CANCER DATASET
```

```
# GMMHMM(With Tuning)[30-70 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =
```

```
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
    '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.GMMHMM(n_components=5,  
random_state=20,covariance_type='diag',algorithm='viterbi',n_iter=10)  
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ("M")
```

```
    else:
```

```
        strings[i] = ("B")
```

```
strings
```

```
from sklearn.metrics import classification_report, confusion_matrix,
accuracy_score
```

```
print("Confusion Matrix:")
print(confusion_matrix(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Performance Evaluation")
print(classification_report(y_test, strings))
```

```
print("-----")
print("-----")
```

```
print("Accuracy:")
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
import seaborn as sns
cm = confusion_matrix(y_test, strings)
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
plt.show()
```

```
#####
#####
#####
#####
```

```
import pandas as pd
import numpy as np
```

Dataset Preparation

```
df = pd.read_csv('wdbc.data',header=None)
```

```
col_name =
```

```
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    , '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
```

```
train_test_split(X,y,train_size=0.7,test_size=0.3,random_state=10)
```

Feature Scaling

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

Classification

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,
```

```
random_state=15,n_iter=10,algorithm='viterbi',params='ste')
```

```
import math
row = len(X_train)
col = len(X_train[0])
new
for i in range(row):
    for j in range(col):
        X_train[i][j] = X_train[i][j]*10
        X_train[i][j] = math.floor(X_train[i][j])
    x = X_train[i].astype(np.int)
    new = np.vstack([new,x])

y = new
y = np.absolute(y)
X_train = y
```

```
import math
row = len(X_test)
col = len(X_test[0])
new
for i in range(row):
    for j in range(col):
        X_test[i][j] = X_test[i][j]*10
        X_test[i][j] = math.floor(X_test[i][j])
    x = X_test[i].astype(np.int)
    new = np.vstack([new,x])
```

```
y = new
y = np.absolute(y)
X_test = y
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ('M')
```

```
    else:
```

```
        strings[i] = ('B')
```

```
strings
```

```
strings = strings[0:171]
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
cm = confusion_matrix(y_test, strings)
```

```
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
```

```
plt.show()
```

```
# BREAST CANCER DATASET
```

```
# MultinomialHMM(With Tuning)[60-40 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =
```

```
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',  
 '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
```

```
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```



```
X_train, X_test, y_train, y_test =  
train_test_split(X,y,train_size=0.6,test_size=0.4,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
```

```
X_train = sc.fit_transform(X_train)
```

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,  
random_state=15,n_iter=10,algorithm='viterbi',params='ste')
```

```
import math
```

```
row = len(X_train)
```

```
col = len(X_train[0])
```

```
new
```

```
for i in range(row):
```

```
    for j in range(col):
```

```
        X_train[i][j] = X_train[i][j]*10
```

```
        X_train[i][j] = math.floor(X_train[i][j])
```

```
        x = X_train[i].astype(np.int)
```

```
        new = np.vstack([new,x])
```

```
y = new
```

```
y = np.absolute(y)
```

```
X_train = y
```

```
import math
row = len(X_test)
col = len(X_test[0])
new
for i in range(row):
    for j in range(col):
        X_test[i][j] = X_test[i][j]*10
        X_test[i][j] = math.floor(X_test[i][j])
        x = X_test[i].astype(np.int)
        new = np.vstack([new,x])
```

```
y = new
y = np.absolute(y)
X_test = y
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ("M")
    else:
        strings[i] = ("B")
```

```
strings
```

```
strings = strings[0:228]
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# BREAST CANCER DATASET  
# MultinomialHMM(With Tuning)[50-50 split]
```

```
import pandas as pd
import numpy as np

# Dataset Preparation
df = pd.read_csv('wdbc.data',header=None)

col_name =
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'
    ,'20','21','22','23','24','25','26','27','28','29','30','31','32']

df.columns = col_name

X = df.drop(['1','Class'], axis=1)
y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.5,test_size=0.5,random_state=10)

# Feature Scaling
from sklearn.preprocessing import StandardScaler

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

# Classification
# from hmmlearn import hmm
```

```
import hmmlearn
classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,
random_state=15,n_iter=10,algorithm='viterbi',params='ste')
```

```
import math
row = len(X_train)
col = len(X_train[0])
new
for i in range(row):
    for j in range(col):
        X_train[i][j] = X_train[i][j]*10
        X_train[i][j] = math.floor(X_train[i][j])
        x = X_train[i].astype(np.int)
        new = np.vstack([new,x])
```

```
y = new
y = np.absolute(y)
X_train = y
```

```
import math
row = len(X_test)
col = len(X_test[0])
new
for i in range(row):
    for j in range(col):
        X_test[i][j] = X_test[i][j]*10
        X_test[i][j] = math.floor(X_test[i][j])
        x = X_test[i].astype(np.int)
        new = np.vstack([new,x])
```

```
y = new
```

```
y = np.absolute(y)
```

```
X_test = y
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
```

```
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
```

```
    if y_pred[i] == 1:
```

```
        strings[i] = ('M')
```

```
    else:
```

```
        strings[i] = ('B')
```

```
strings
```

```
strings = strings[0:285]
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")
```

```
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")
```

```
print(classification_report(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Accuracy:")
```

```
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
cm = confusion_matrix(y_test, strings)
```

```
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
```

```
plt.show()
```

```
# BREAST CANCER DATASET
```

```
# MultinomialHMM(With Tuning)[40-60 split]
```

```
import pandas as pd
```

```
import numpy as np
```

```
# Dataset Preparation
```

```
df = pd.read_csv('wdbc.data', header=None)
```

```
col_name =
```

```
['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19'  
    '20','21','22','23','24','25','26','27','28','29','30','31','32']
```

```
df.columns = col_name
```

```
X = df.drop(['1','Class'], axis=1)
y = df['Class']
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test =
train_test_split(X,y,train_size=0.4,test_size=0.6,random_state=10)
```

```
# Feature Scaling
```

```
from sklearn.preprocessing import StandardScaler
```

```
sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
import hmmlearn
classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,
random_state=15,n_iter=10,algorithm='viterbi',params='ste')
```

```
import math
row = len(X_train)
col = len(X_train[0])
new
for i in range(row):
    for j in range(col):
        X_train[i][j] = X_train[i][j]*10
        X_train[i][j] = math.floor(X_train[i][j])
        x = X_train[i].astype(np.int)
        new = np.vstack([new,x])
```



```
y = new
y = np.absolute(y)
X_train = y
```

```
import math
row = len(X_test)
col = len(X_test[0])
new
for i in range(row):
    for j in range(col):
        X_test[i][j] = X_test[i][j]*10
        X_test[i][j] = math.floor(X_test[i][j])
        x = X_test[i].astype(np.int)
        new = np.vstack([new,x])
```

```
y = new
y = np.absolute(y)
X_test = y
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):
    if y_pred[i] == 1:
        strings[i] = ("M")
```

```
else:  
    strings[i] = ("B")
```

```
strings  
strings = strings[0:342]
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

BREAST CANCER DATASET

MultinomialHMM(With Tuning)[30-70 split]

import pandas as pd

import numpy as np

Dataset Preparation

df = pd.read_csv('wdbc.data',header=None)

col_name =

**['1','Class','3','4','5','6','7','8','9','10','11','12','13','14','15','16','17','18','19',
'20','21','22','23','24','25','26','27','28','29','30','31','32']**

df.columns = col_name

X = df.drop(['1','Class'], axis=1)

y = df['Class']

from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test =

train_test_split(X,y,train_size=0.3,test_size=0.7,random_state=10)

Feature Scaling

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

X_train = sc.fit_transform(X_train)

```
X_test = sc.transform(X_test)
```

```
# Classification
```

```
# from hmmlearn import hmm
```

```
import hmmlearn
```

```
classifier = hmmlearn.hmm.MultinomialHMM(n_components=4,  
random_state=15,n_iter=10,algorithm='viterbi',params='ste')
```

```
import math
```

```
row = len(X_train)
```

```
col = len(X_train[0])
```

```
new
```

```
for i in range(row):
```

```
    for j in range(col):
```

```
        X_train[i][j] = X_train[i][j]*10
```

```
        X_train[i][j] = math.floor(X_train[i][j])
```

```
        x = X_train[i].astype(np.int)
```

```
        new = np.vstack([new,x])
```

```
y = new
```

```
y = np.absolute(y)
```

```
X_train = y
```

```
import math
```

```
row = len(X_test)
```

```
col = len(X_test[0])
```

```
new
```

```
for i in range(row):
```

```
    for j in range(col):
```

```
        X_test[i][j] = X_test[i][j]*10
```

```
        X_test[i][j] = math.floor(X_test[i][j])
```

```
x = X_test[i].astype(np.int)  
new = np.vstack([new,x])
```

```
y = new  
y = np.absolute(y)  
X_test = y
```

```
classifier.fit(X_train)
```

```
y_pred = classifier.predict(X_test)
```

```
size = len(y_pred)  
strings = np.empty(size, np.unicode_)
```

```
for i in range (size):  
    if y_pred[i] == 1:  
        strings[i] = ("M")  
    else:  
        strings[i] = ("B")
```

```
strings  
strings = strings[0:399]
```

```
from sklearn.metrics import classification_report, confusion_matrix,  
accuracy_score
```

```
print("Confusion Matrix:")  
print(confusion_matrix(y_test, strings))
```

```
print("-----")
```

```
print("-----")
```

```
print("Performance Evaluation")  
print(classification_report(y_test, strings))
```

```
print("-----")  
print("-----")
```

```
print("Accuracy:")  
print(accuracy_score(y_test, strings))
```

```
import matplotlib.pyplot as plt  
import seaborn as sns  
cm = confusion_matrix(y_test, strings)  
sns.heatmap(cm, annot=True, fmt="d", cmap='Blues')  
plt.show()
```

```
# CIFAR 10 DATASET
```

```
import pandas as pd  
import numpy as np  
import matplotlib.pyplot as plt  
import tensorflow as tf
```

```
# dataset preparation
```

```
from tensorflow.keras import datasets, layers, models
```

```
(train_images, train_labels), (test_images, test_labels) =  
datasets.cifar10.load_data()
```

```
# Normalize pixel values to be within 0 , 1
train_images , test_images = train_images/255.0 , test_images/255.0

input_shape = train_images[0].shape

model = models.Sequential()
model.add(layers.Conv2D(32,(3,3),activation='relu',input_shape=input_shape))
model.add(layers.MaxPool2D(2,2))
model.add(layers.Conv2D(64,(3,3),activation='relu'))
model.add(layers.MaxPool2D(2,2))
model.add(layers.Conv2D(64,(3,3),activation='relu'))

model.add(layers.Flatten())
model.add(layers.Dense(64,activation='relu'))
model.add(layers.Dense(10))

model.summary()

model.compile(optimizer='adam',loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),metrics=['accuracy'])

history =
model.fit(train_images,train_labels,epochs=20,validation_data=(test_images,test_labels))

plt.plot(history.history['accuracy'],label='accuracy')
plt.plot(history.history['val_accuracy'],label='val_accuracy')
plt.xlabel('epoch')
plt.ylabel('accuracy')
plt.ylim([0.5,1])
```

```
plt.legend(loc='lower right')
```

```
plt.show()
```

```
test_loss , test_acc = model.evaluate(test_images,test_labels,verbose=2)
```

```
# MNIST DATASET
```

```
import pandas as pd
```

```
import numpy as np
```

```
import matplotlib.pyplot as plt
```

```
import tensorflow as tf
```

```
# dataset preparation
```

```
from tensorflow.keras import datasets,layers,models
```

```
(train_images,train_labels) , (test_images,test_labels) = datasets.mnist.load_data()
```

```
# Normalize pixel values to be within 0 , 1
```

```
train_images , test_images = train_images/255.0 , test_images/255.0
```

```
train_images = np.reshape(train_images, train_images.shape + (1,))
```

```
test_images = np.reshape(test_images, test_images.shape + (1,))
```

```
train_images[0].shape
```

```
model = models.Sequential()
```

```
model.add(layers.Conv2D(32,(3,3),activation='relu',input_shape=(28,28,1)))
```

```
model.add(layers.MaxPool2D(2,2))
```

```
model.add(layers.Conv2D(64,(3,3),activation='relu'))
```

```
model.add(layers.MaxPool2D(2,2))
```



```
model.add(layers.Conv2D(64,(3,3),activation='relu'))
```

```
model.add(layers.Flatten())
```

```
model.add(layers.Dense(64,activation='relu'))
```

```
model.add(layers.Dense(10))
```

```
model.summary()
```

```
model.compile(optimizer='adam',loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),metrics=['accuracy'])
```

```
history =
```

```
model.fit(train_images,train_labels,epochs=20,validation_data=(test_images,test_labels))
```

```
plt.plot(history.history['accuracy'],label='accuracy')
```

```
plt.plot(history.history['val_accuracy'],label='val_accuracy')
```

```
plt.xlabel('epoch')
```

```
plt.ylabel('accuracy')
```

```
plt.ylim([0.5,1])
```

```
plt.legend(loc='lower right')
```

```
plt.show()
```

```
test_loss , test_acc = model.evaluate(test_images,test_labels,verbose=2)
```

```
# SAVEE DATASET
```

```
!unzip "/content/drive/MyDrive/SaveeDataset.zip"
```

```
import librosa
```

```

import numpy as np

input_length = 16000*5

batch_size = 32

n_mels = 320

def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,
#log_specgram
    step_size=10, eps=1e-10):

    mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,
n_mels= n_mels)
    mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40

    return mel_db.T

def load_audio_file(file_path, input_length=input_length):
    data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000
    if len(data)>input_length:
        max_offset = len(data)-input_length

        offset = np.random.randint(max_offset)

        data = data[offset:(input_length+offset)]

    else:
        if input_length > len(data):
            max_offset = input_length - len(data)

```

```
offset = np.random.randint(max_offset)
else:
offset = 0
data = np.pad(data, (offset, input_length - len(data) - offset), "constant")
```

```
data = preprocess_audio_mel_T(data)
return data
```

Preprocessing the dataset

```
import os
from scipy.io import wavfile
import librosa
import matplotlib.pyplot as plt
import numpy as np
```

```
rootDirectory = "/content/AudioData/"
personNames = ["DC","JE","JK","KL"]
```

```
classes = ["a" , "d" , "f" , "h" , "n" , "sa" , "su" ]
```

```
X = list()
y = list()
```

```
for person in personNames:
    directory = os.path.join(rootDirectory,person)
    for filename in os.listdir(directory):
        filePath = os.path.join(directory, filename)
        data = load_audio_file(file_path=filePath)
        data = np.reshape(data, data.shape + (1,))
        if(filename[0:1] in classes):
            X.append(data)
            y.append(classes.index(filename[0:1]))
```

```
elif(filename[0:2] in classes):  
    X.append(data)  
    y.append(classes.index(filename[0:2]))
```

```
X = np.asarray(X, dtype=np.float32)  
y = np.asarray(y, dtype=np.float32)
```

```
import pandas as pd  
import numpy as np  
import matplotlib.pyplot as plt  
import tensorflow as tf
```

```
# dataset preparation
```

```
from tensorflow.keras import datasets, layers, models  
from sklearn.model_selection import train_test_split  
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, train_size= 0.7  
, random_state=10)
```

```
model = models.Sequential()  
model.add(layers.Conv2D(32,(3,3),activation='relu',input_shape=(157,320,1)))  
model.add(layers.MaxPool2D(2,2))  
model.add(layers.Conv2D(64,(3,3),activation='relu'))  
model.add(layers.MaxPool2D(2,2))  
model.add(layers.Conv2D(64,(3,3),activation='relu'))
```

```
model.add(layers.Flatten())  
model.add(layers.Dense(64,activation='relu'))  
model.add(layers.Dense(10))
```

```
model.summary()
```

```
model.compile(optimizer='adam',loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),metrics=['accuracy'])
```

```
history = model.fit(X_train,y_train,epochs=30,validation_data=(X_test,y_test))
```

```
plt.plot(history.history['accuracy'],label='accuracy')  
plt.plot(history.history['val_accuracy'],label='val_accuracy')  
plt.xlabel('epoch')  
plt.ylabel('accuracy')  
plt.ylim([0.5,1])  
plt.legend(loc='lower right')
```

```
plt.show()
```

```
test_loss , test_acc = model.evaluate(X_test,y_test,verbose=2)
```

```
# EmoDB DATASET
```

```
!unzip "/content/drive/MyDrive/EmoDB.zip"
```

```
import librosa  
import numpy as np
```

```
input_length = 16000*5
```

```
batch_size = 32
```

```
n_mels = 320
```

```
def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,  
#log_specgram
```

step_size=10, eps=1e-10):

**mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,
n_mels= n_mels)**

mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40

return mel_db.T

def load_audio_file(file_path, input_length=input_length):

data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000

if len(data)>input_length:

max_offset = len(data)-input_length

offset = np.random.randint(max_offset)

data = data[offset:(input_length+offset)]

else:

if input_length > len(data):

max_offset = input_length - len(data)

offset = np.random.randint(max_offset)

else:

offset = 0

data = np.pad(data, (offset, input_length - len(data) - offset), "constant")

data = preprocess_audio_mel_T(data)

return data

Preprocessing the dataset

import os

```
from scipy.io import wavfile
import librosa
import matplotlib.pyplot as plt
import numpy as np

directory = "/content/wav/"

classes = ["W" ,"L" ,"E" ,"A" , "F" ,"T" ,"N" ]

X = list()
y = list()

for filename in os.listdir(directory):
    filePath = os.path.join(directory, filename)
    data = load_audio_file(file_path=filePath)
    data = np.reshape(data, data.shape + (1,))
    if(filename[5:6] in classes):
        X.append(data)
        y.append(classes.index(filename[5:6]))

X = np.asarray(X, dtype=np.float32)
y = np.asarray(y, dtype=np.float32)

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import tensorflow as tf

# dataset preparation

from tensorflow.keras import datasets, layers, models
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, train_size= 0.7,random_state=10)
```

```
model = models.Sequential()  
model.add(layers.Conv2D(32,(3,3),activation='relu',input_shape=(157,320,1)))  
model.add(layers.MaxPool2D(2,2))  
model.add(layers.Conv2D(64,(3,3),activation='relu'))  
model.add(layers.MaxPool2D(2,2))  
model.add(layers.Conv2D(64,(3,3),activation='relu'))
```

```
model.add(layers.Flatten())  
model.add(layers.Dense(64,activation='relu'))  
model.add(layers.Dense(10))
```

```
model.summary()
```

```
model.compile(optimizer='adam',loss=tf.keras.losses.SparseCategoricalCrossentropy(from_logits=True),metrics=['accuracy'])
```

```
history = model.fit(X_train,y_train,epochs=20,validation_data=(X_test,y_test))
```

```
plt.plot(history.history['accuracy'],label='accuracy')  
plt.plot(history.history['val_accuracy'],label='val_accuracy')  
plt.xlabel('epoch')  
plt.ylabel('accuracy')  
plt.ylim([0,1])  
plt.legend(loc='lower right')
```

```
plt.show()
```

```
test_loss , test_acc = model.evaluate(X_test,y_test,verbose=2)
```


#@title

```
from google.colab import drive  
drive.mount('/content/drive')
```

#@title

```
import tensorflow as tf  
from tensorflow import keras  
import matplotlib.pyplot as plt  
%matplotlib inline  
import numpy as np  
import skimage.transform  
from __future__ import print_function
```

```
!pip install keras_applications
```

```
import numpy as np  
import warnings
```

```
from keras.models import Model  
from keras.layers import Flatten  
from keras.layers import Dense  
from keras.layers import Input  
from keras.layers import Conv2D  
from keras.layers import MaxPooling2D  
from keras.layers import GlobalMaxPooling2D  
from keras.layers import GlobalAveragePooling2D  
from keras.preprocessing import image  
from keras.utils import layer_utils  
from keras.utils.data_utils import get_file  
from keras import backend as K  
from keras.applications.imagenet_utils import decode_predictions  
from keras.applications.imagenet_utils import preprocess_input
```

```
from keras_applications.imagenet_utils import _obtain_input_shape  
from keras.utils.layer_utils import get_source_inputs
```

```
#@title
```

```
def load_preprocess_training_batch(X_train):
```

```
    new = []
```

```
    for item in X_train:
```

```
        tmpFeature = skimage.transform.resize(item, (224, 224), mode='constant')
```

```
        new.append(tmpFeature)
```

```
    return new
```

```
#@title
```

```
def preprocess_data(X_train):
```

```
    for item in X_train:
```

```
        item = np.expand_dims(item, axis=0)
```

```
        item = preprocess_input(item)
```

```
    return X_train
```

```
#@title
```

```
WEIGHTS_PATH = 'https://github.com/fchollet/deep-learning-
```

```
models/releases/download/v0.1/vgg16_weights_tf_dim_ordering_tf_kernels.h5'
```

```
WEIGHTS_PATH_NO_TOP = 'https://github.com/fchollet/deep-learning-
```

```
models/releases/download/v0.1/vgg16_weights_tf_dim_ordering_tf_kernels_notop.  
h5'
```

```
def VGG16(include_top=True, weights='imagenet',  
          input_tensor=None, input_shape=None,  
          pooling=None,  
          classes=1000):  
    """Instantiates the VGG16 architecture.
```

Optionally loads weights pre-trained on ImageNet. Note that when using TensorFlow, for best performance you should set ``image_data_format="channels_last"`` in your Keras config at `~/.keras/keras.json`.

The model and the weights are compatible with both TensorFlow and Theano. The data format convention used by the model is the one specified in your Keras config file.

Arguments

include_top: whether to include the 3 fully-connected layers at the top of the network.

weights: one of ``None`` (random initialization) or `"imagenet"` (pre-training on ImageNet).

input_tensor: optional Keras tensor (i.e. output of ``layers.Input()``) to use as image input for the model.

input_shape: optional shape tuple, only to be specified if ``include_top`` is `False` (otherwise the input shape has to be ``(224, 224, 3)`` (with ``channels_last`` data format) or ``(3, 224, 244)`` (with ``channels_first`` data format).

It should have exactly 3 inputs channels, and width and height should be no smaller than 48.

E.g. ``(200, 200, 3)`` would be one valid value.

pooling: Optional pooling mode for feature extraction when ``include_top`` is ``False``.

- ``None`` means that the output of the model will be the 4D tensor output of the last convolutional layer.
- ``avg`` means that global average pooling will be applied to the output of the last convolutional layer, and thus the output of the model will be a 2D tensor.
- ``max`` means that global max pooling will be applied.

classes: optional number of classes to classify images into, only to be specified if ``include_top`` is `True`, and if no ``weights`` argument is specified.

Returns

A Keras model instance.

Raises

ValueError: in case of invalid argument for ``weights``, or invalid input shape.

"""

if `weights` not in `{'imagenet', None}`:

raise ValueError('The ``weights`` argument should be either '
 ``None`` (random initialization) or ``imagenet`` '
 '(pre-training on ImageNet).')

if `weights == 'imagenet'` and `include_top` and `classes != 1000`:

raise ValueError('If using ``weights`` as imagenet with ``include_top``
 ' as true, ``classes`` should be 1000')

Determine proper input shape

`input_shape = _obtain_input_shape(input_shape,`

```
default_size=224,  
min_size=48,  
data_format=K.image_data_format(),  
require_flatten=include_top)
```

```
if input_tensor is None:
```

```
img_input = Input(shape=input_shape)
```

```
else:
```

```
if not K.is_keras_tensor(input_tensor):
```

```
img_input = Input(tensor=input_tensor, shape=input_shape)
```

```
else:
```

```
img_input = input_tensor
```

```
# Block 1
```

```
x = Conv2D(64, (3, 3), activation='relu', padding='same',  
name='block1_conv1')(img_input)
```

```
x = Conv2D(64, (3, 3), activation='relu', padding='same',  
name='block1_conv2')(x)
```

```
x = MaxPooling2D((2, 2), strides=(2, 2), name='block1_pool')(x)
```

```
# Block 2
```

```
x = Conv2D(128, (3, 3), activation='relu', padding='same',  
name='block2_conv1')(x)
```

```
x = Conv2D(128, (3, 3), activation='relu', padding='same',  
name='block2_conv2')(x)
```

```
x = MaxPooling2D((2, 2), strides=(2, 2), name='block2_pool')(x)
```

```
# Block 3
```

```
x = Conv2D(256, (3, 3), activation='relu', padding='same',  
name='block3_conv1')(x)
```

```
x = Conv2D(256, (3, 3), activation='relu', padding='same',  
name='block3_conv2')(x)
```

```
x = Conv2D(256, (3, 3), activation='relu', padding='same',  
name='block3_conv3')(x)  
x = MaxPooling2D((2, 2), strides=(2, 2), name='block3_pool')(x)
```

```
# Block 4
```

```
x = Conv2D(512, (3, 3), activation='relu', padding='same',  
name='block4_conv1')(x)  
x = Conv2D(512, (3, 3), activation='relu', padding='same',  
name='block4_conv2')(x)  
x = Conv2D(512, (3, 3), activation='relu', padding='same',  
name='block4_conv3')(x)  
x = MaxPooling2D((2, 2), strides=(2, 2), name='block4_pool')(x)
```

```
# Block 5
```

```
x = Conv2D(512, (3, 3), activation='relu', padding='same',  
name='block5_conv1')(x)  
x = Conv2D(512, (3, 3), activation='relu', padding='same',  
name='block5_conv2')(x)  
x = Conv2D(512, (3, 3), activation='relu', padding='same',  
name='block5_conv3')(x)  
x = MaxPooling2D((2, 2), strides=(2, 2), name='block5_pool')(x)
```

```
if include_top:
```

```
# Classification block
```

```
x = Flatten(name='flatten')(x)  
x = Dense(4096, activation='relu', name='fc1')(x)  
x = Dense(4096, activation='relu', name='fc2')(x)  
x = Dense(classes, activation='softmax', name='predictions')(x)  
else:  
if pooling == 'avg':  
x = GlobalAveragePooling2D()(x)  
elif pooling == 'max':
```

```
x = GlobalMaxPooling2D()(x)
```

```
# Ensure that the model takes into account  
# any potential predecessors of `input_tensor`.
```

```
if input_tensor is not None:
```

```
inputs = get_source_inputs(input_tensor)
```

```
else:
```

```
inputs = img_input
```

```
# Create model.
```

```
model = Model(inputs, x, name='vgg16')
```

```
# load weights
```

```
if weights == 'imagenet':
```

```
if include_top:
```

```
weights_path = get_file('vgg16_weights_tf_dim_ordering_tf_kernels.h5',  
                        WEIGHTS_PATH,  
                        cache_subdir='models')
```

```
else:
```

```
weights_path =
```

```
get_file('vgg16_weights_tf_dim_ordering_tf_kernels_notop.h5',  
        WEIGHTS_PATH_NO_TOP,  
        cache_subdir='models')
```

```
model.load_weights(weights_path)
```

```
if K.backend() == 'theano':
```

```
layer_utils.convert_all_kernels_in_model(model)
```

```
if K.image_data_format() == 'channels_first':
```

```
if include_top:
```

```
    maxpool = model.get_layer(name='block5_pool')
```

```
    shape = maxpool.output_shape[1:]
```

```
    dense = model.get_layer(name='fc1')
```

```
layer_utils.convert_dense_weights_data_format(dense, shape,
'channels_first')
```

```
if K.backend() == 'tensorflow':
    warnings.warn('You are using the TensorFlow backend, yet you '
        'are using the Theano '
        'image data format convention '
        '('image_data_format="channels_first"'). '
        'For best performance, set '
        '`image_data_format="channels_last"` in '
        'your Keras config '
        'at ~/.keras/keras.json.')
return model
```

```
#@title
```

```
import tensorflow as tf
from tensorflow import keras
import matplotlib.pyplot as plt
%matplotlib inline
import numpy as np
import skimage.transform
```

```
# CIFAR-10 Dataset
```

```
#@title
```

```
(X_train, y_train) , (X_test, y_test) = keras.datasets.cifar10.load_data()
```

```
X_train = X_train[0:2000]
y_train = y_train[0:2000]
X_test = X_test[0:2000]
y_test = y_test[0:2000]
```


#@title

X_train_resized = load_preprocess_training_batch(X_train)

X_test_resized = load_preprocess_training_batch(X_test)

#@title

X_train_resized = np.array(X_train_resized)

X_test_resized = np.array(X_test_resized)

#@title

X_train_resized = X_train_resized / 255

X_test_resized = X_test_resized / 255

#@title

X_train_resized = preprocess_data(X_train_resized)

X_test_resized = preprocess_data(X_test_resized)

#@title

model = VGG16(include_top=True, weights='imagenet')

**model.compile(optimizer='SGD',
 loss='sparse_categorical_crossentropy',
 metrics=['accuracy'])**

history = model.fit(X_train_resized, y_train, epochs=5)

img_path = 'a.jpg'

img = image.load_img(img_path, target_size=(224, 224))

x = image.img_to_array(img)

x = np.expand_dims(x, axis=0)

x = preprocess_input(x)

print('Input image shape:', x.shape)

```
# preds = model.predict(x)
# print('Predicted:', decode_predictions(preds))
```

```
#@title
model.evaluate(X_test_resized, y_test)
```

```
# MNIST Dataset
```

```
#@title
(X_train, y_train) , (X_test, y_test) = keras.datasets.mnist.load_data()
```

```
X_train = X_train[0:2000]
y_train = y_train[0:2000]
X_test = X_test[0:2000]
y_test = y_test[0:2000]
```

```
#@title
X_train_resized = load_preprocess_training_batch(X_train)
X_test_resized = load_preprocess_training_batch(X_test)
```

```
X_train_resized = np.array(X_train_resized)
X_test_resized = np.array(X_test_resized)
```

```
X_train_resized = X_train_resized / 255.0
X_test_resized = X_test_resized / 255.0
```

```
X_train_resized = preprocess_data(X_train_resized)
X_test_resized = preprocess_data(X_test_resized)
```

```
#@title
import cv2
```

```
X_train_new = list()
```

```
for i in range(len(X_train_resized)):  
    g = X_train_resized[i]  
    X_train_new.append(cv2.merge([g,g,g]))
```

```
X_train_new = np.asarray(X_train_new,dtype=np.float32)
```

```
X_test_new = list()
```

```
for i in range(len(X_test_resized)):  
    g = X_test_resized[i]  
    X_test_new.append(cv2.merge([g,g,g]))
```

```
X_test_new = np.asarray(X_test_new,dtype=np.float32)
```

```
#@title
```

```
model = VGG16(include_top=True, weights='imagenet')
```

```
model.compile(optimizer='SGD',  
              loss='sparse_categorical_crossentropy',  
              metrics=['accuracy'])
```

```
history = model.fit(X_train_new, y_train, epochs=5)
```

```
#@title
```

```
model.evaluate(X_test_new, y_test)
```

```
# SAVEE Dataset
```

```
#@title
```

```
!unzip "/content/drive/MyDrive/SaveeDataset.zip"
```

```
#@title
```

```
import librosa
```

```
import numpy as np
```

```
input_length = 16000*5
```

```
batch_size = 32
```

```
n_mels = 320
```

```
def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,
```

```
#log_specgram
```

```
    step_size=10, eps=1e-10):
```

```
        mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,  
n_mels= n_mels)
```

```
        mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40
```

```
        return mel_db.T
```

```
def load_audio_file(file_path, input_length=input_length):
```

```
    data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000
```

```
    if len(data)>input_length:
```

```
        max_offset = len(data)-input_length
```

```
        offset = np.random.randint(max_offset)
```

```
        data = data[offset:(input_length+offset)]
```

```
else:
    if input_length > len(data):
        max_offset = input_length - len(data)

        offset = np.random.randint(max_offset)
    else:
        offset = 0
    data = np.pad(data, (offset, input_length - len(data) - offset), "constant")

data = preprocess_audio_mel_T(data)
return data
```

```
# @title
# Preprocessing the dataset
import os
from scipy.io import wavfile
import librosa
import matplotlib.pyplot as plt
import numpy as np
import cv2

rootDirectory = "/content/AudioData/"
personNames = ["DC", "JE", "JK", "KL"]

classes = ["a", "d", "f", "h", "n", "sa", "su"]

X = list()
y = list()

for person in personNames:
    directory = os.path.join(rootDirectory, person)
    for filename in os.listdir(directory):
```

```
filePath = os.path.join(directory, filename)
a = load_audio_file(file_path=filePath)
data = cv2.merge([a,a,a])
# data = np.reshape(data, data.shape + (1,))
if(filename[0:1] in classes):
X.append(data)
y.append(classes.index(filename[0:1]))
elif(filename[0:2] in classes):
X.append(data)
y.append(classes.index(filename[0:2]))
```

#@title

```
X = np.asarray(X, dtype=np.float32)
y = np.asarray(y, dtype=np.float32)
```

#@title

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import tensorflow as tf
```

dataset preparation

```
from tensorflow.keras import datasets, layers, models
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.5, train_size= 0.5
,random_state=10)
```

#@title

```
X_train_resized = load_preprocess_training_batch(X_train)
X_test_resized = load_preprocess_training_batch(X_test)
```

```
X_train_resized = np.array(X_train_resized)
```

```
X_test_resized = np.array(X_test_resized)
```

```
X_train_resized = preprocess_data(X_train_resized)
```

```
X_test_resized = preprocess_data(X_test_resized)
```

```
#@title
```

```
model = VGG16(include_top=True, weights='imagenet')
```

```
model.compile(optimizer='SGD',  
              loss='sparse_categorical_crossentropy',  
              metrics=['accuracy'])
```

```
history = model.fit(X_train_resized, y_train, epochs=50)
```

```
#@title
```

```
model.evaluate(X_test_resized, y_test)
```

```
# EmoDb Dataset
```

```
#@title
```

```
!unzip "/content/drive/MyDrive/EmoDB.zip"
```

```
#@title
```

```
import librosa
```

```
import numpy as np
```

```
input_length = 16000*5
```

```
batch_size = 32
```

```
n_mels = 320
```

```

def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,
#log_specgram
    step_size=10, eps=1e-10):

    mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,
n_mels= n_mels)
    mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40

    return mel_db.T

def load_audio_file(file_path, input_length=input_length):
    data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000
    if len(data)>input_length:
        max_offset = len(data)-input_length

        offset = np.random.randint(max_offset)

        data = data[offset:(input_length+offset)]

    else:
        if input_length > len(data):
            max_offset = input_length - len(data)

            offset = np.random.randint(max_offset)
            else:
                offset = 0
            data = np.pad(data, (offset, input_length - len(data) - offset), "constant")

    data = preprocess_audio_mel_T(data)
    return data

```



```
#@title  
# Preprocessing the dataset  
import os  
from scipy.io import wavfile  
import librosa  
import matplotlib.pyplot as plt  
import numpy as np  
import cv2  
  
directory = "/content/wav/"  
  
classes = ["W" , "L" , "E" , "A" , "F" , "T" , "N" ]  
  
X = list()  
y = list()  
  
for filename in os.listdir(directory):  
    filePath = os.path.join(directory, filename)  
    a = load_audio_file(file_path=filePath)  
    data = cv2.merge([a,a,a])  
    if(filename[5:6] in classes):  
        X.append(data)  
        y.append(classes.index(filename[5:6]))  
  
#@title  
X = np.asarray(X, dtype=np.float32)  
y = np.asarray(y, dtype=np.float32)  
  
#@title  
import pandas as pd  
import numpy as np
```

```
import matplotlib.pyplot as plt
import tensorflow as tf

# dataset preparation

from tensorflow.keras import datasets, layers, models
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.5, train_size= 0.5
,random_state=10)

#@title
X_train_resized = load_preprocess_training_batch(X_train)
X_test_resized = load_preprocess_training_batch(X_test)

X_train_resized = np.array(X_train_resized)
X_test_resized = np.array(X_test_resized)

X_train_resized = preprocess_data(X_train_resized)
X_test_resized = preprocess_data(X_test_resized)

#@title
model = VGG16(include_top=True, weights='imagenet')

model.compile(optimizer='SGD',
              loss='sparse_categorical_crossentropy',
              metrics=['accuracy'])

history = model.fit(X_train_resized, y_train, epochs=20)

#@title
model.evaluate(X_test_resized, y_test)
from google.colab import drive
```

```
drive.mount('/content/drive')
```

```
from __future__ import print_function
```

```
import numpy as np
```

```
import warnings
```

```
!pip install keras_applications
```

```
from keras.layers import Input
```

```
from keras import layers
```

```
from keras.layers import Dense
```

```
from keras.layers import Activation
```

```
from keras.layers import Flatten
```

```
from keras.layers import Conv2D
```

```
from keras.layers import MaxPooling2D
```

```
from keras.layers import GlobalMaxPooling2D
```

```
from keras.layers import ZeroPadding2D
```

```
from keras.layers import AveragePooling2D
```

```
from keras.layers import GlobalAveragePooling2D
```

```
from keras.layers import BatchNormalization
```

```
from keras.models import Model
```

```
from keras.preprocessing import image
```

```
import keras.backend as K
```

```
from keras.utils import layer_utils
```

```
from keras.utils.data_utils import get_file
```

```
from keras.applications.imagenet_utils import decode_predictions
```

```
from keras.applications.imagenet_utils import preprocess_input
```

```
from keras_applications.imagenet_utils import _obtain_input_shape
```

```
from keras.utils.layer_utils import get_source_inputs
```

```
import tensorflow as tf
```

```
from tensorflow import keras  
import matplotlib.pyplot as plt  
%matplotlib inline  
import numpy as np  
import skimage.transform
```

```
WEIGHTS_PATH = 'https://github.com/fchollet/deep-learning-  
models/releases/download/v0.2/resnet50_weights_tf_dim_ordering_tf_kernels.h5'  
WEIGHTS_PATH_NO_TOP = 'https://github.com/fchollet/deep-learning-  
models/releases/download/v0.2/resnet50_weights_tf_dim_ordering_tf_kernels_not  
op.h5'
```

```
def identity_block(input_tensor, kernel_size, filters, stage, block):  
    """The identity block is the block that has no conv layer at shortcut.  
    # Arguments  
    input_tensor: input tensor  
    kernel_size: default 3, the kernel size of middle conv layer at main path  
    filters: list of integers, the filters of 3 conv layer at main path  
    stage: integer, current stage label, used for generating layer names  
    block: 'a','b'..., current block label, used for generating layer names  
    # Returns  
    Output tensor for the block.  
    """  
  
    filters1, filters2, filters3 = filters  
    if K.image_data_format() == 'channels_last':  
    bn_axis = 3  
    else:  
    bn_axis = 1  
    conv_name_base = 'res' + str(stage) + block + '_branch'  
    bn_name_base = 'bn' + str(stage) + block + '_branch'
```

```

x = Conv2D(filters1, (1, 1), name=conv_name_base + '2a')(input_tensor)
x = BatchNormalization(axis=bn_axis, name=bn_name_base + '2a')(x)
x = Activation('relu')(x)

```

```

x = Conv2D(filters2, kernel_size,
           padding='same', name=conv_name_base + '2b')(x)
x = BatchNormalization(axis=bn_axis, name=bn_name_base + '2b')(x)
x = Activation('relu')(x)

```

```

x = Conv2D(filters3, (1, 1), name=conv_name_base + '2c')(x)
x = BatchNormalization(axis=bn_axis, name=bn_name_base + '2c')(x)

```

```

x = layers.add([x, input_tensor])
x = Activation('relu')(x)
return x

```

```

def conv_block(input_tensor, kernel_size, filters, stage, block, strides=(2, 2)):
    """conv_block is the block that has a conv layer at shortcut
    # Arguments
    input_tensor: input tensor
    kernel_size: default 3, the kernel size of middle conv layer at main path
    filters: list of integers, the filters of 3 conv layer at main path
    stage: integer, current stage label, used for generating layer names
    block: 'a','b'..., current block label, used for generating layer names
    # Returns
    Output tensor for the block.
    Note that from stage 3, the first conv layer at main path is with strides=(2,2)
    And the shortcut should have strides=(2,2) as well
    """
    filters1, filters2, filters3 = filters
    if K.image_data_format() == 'channels_last':

```

```
bn_axis = 3
else:
bn_axis = 1
conv_name_base = 'res' + str(stage) + block + '_branch'
bn_name_base = 'bn' + str(stage) + block + '_branch'
```

```
x = Conv2D(filters1, (1, 1), strides=strides,
           name=conv_name_base + '2a')(input_tensor)
x = BatchNormalization(axis=bn_axis, name=bn_name_base + '2a')(x)
x = Activation('relu')(x)
```

```
x = Conv2D(filters2, kernel_size, padding='same',
           name=conv_name_base + '2b')(x)
x = BatchNormalization(axis=bn_axis, name=bn_name_base + '2b')(x)
x = Activation('relu')(x)
```

```
x = Conv2D(filters3, (1, 1), name=conv_name_base + '2c')(x)
x = BatchNormalization(axis=bn_axis, name=bn_name_base + '2c')(x)
```

```
shortcut = Conv2D(filters3, (1, 1), strides=strides,
                  name=conv_name_base + '1')(input_tensor)
shortcut = BatchNormalization(axis=bn_axis, name=bn_name_base +
'1')(shortcut)
```

```
x = layers.add([x, shortcut])
x = Activation('relu')(x)
return x
```

```
def ResNet50(include_top=True, weights='imagenet',
             input_tensor=None, input_shape=None,
             pooling=None,
```

classes=1000):

"""Instantiates the ResNet50 architecture.

Optionally loads weights pre-trained on ImageNet. Note that when using TensorFlow, for best performance you should set `image_data_format='channels_last'` in your Keras config at ~/.keras/keras.json.

The model and the weights are compatible with both TensorFlow and Theano. The data format convention used by the model is the one specified in your Keras config file.

Arguments

include_top: whether to include the fully-connected layer at the top of the network.

weights: one of `None` (random initialization) or "imagenet" (pre-training on ImageNet).

input_tensor: optional Keras tensor (i.e. output of `layers.Input()`) to use as image input for the model.

input_shape: optional shape tuple, only to be specified if `include_top` is False (otherwise the input shape has to be `(224, 224, 3)` (with `channels_last` data format) or `(3, 224, 244)` (with `channels_first` data format).

It should have exactly 3 inputs channels, and width and height should be no smaller than 197. E.g. `(200, 200, 3)` would be one valid value.

pooling: Optional pooling mode for feature extraction when `include_top` is `False`.

- `None` means that the output of the model will be the 4D tensor output of the last convolutional layer.
- `avg` means that global average pooling will be applied to the output of the

last convolutional layer, and thus
the output of the model will be a 2D tensor.

- ``max`` means that global max pooling will
be applied.

`classes`: optional number of classes to classify images
into, only to be specified if ``include_top`` is True, and
if no ``weights`` argument is specified.

Returns

A Keras model instance.

Raises

ValueError: in case of invalid argument for ``weights``,
or invalid input shape.

"""

if `weights` not in `{'imagenet', None}`:

raise ValueError('The ``weights`` argument should be either '
 '``None`` (random initialization) or ``imagenet`` '
 '(pre-training on ImageNet).')

if `weights == 'imagenet'` and `include_top` and `classes != 1000`:

raise ValueError('If using ``weights`` as imagenet with ``include_top`` '
 'as true, ``classes`` should be 1000')

Determine proper input shape

`input_shape = _obtain_input_shape(input_shape,
 default_size=224,
 min_size=197,
 data_format=K.image_data_format(),
 require_flatten=include_top)`

if `input_tensor` is None:

`img_input = Input(shape=input_shape)`

else:


```

if not K.is_keras_tensor(input_tensor):
    img_input = Input(tensor=input_tensor, shape=input_shape)
else:
    img_input = input_tensor
if K.image_data_format() == 'channels_last':
    bn_axis = 3
else:
    bn_axis = 1

x = ZeroPadding2D((3, 3))(img_input)
x = Conv2D(64, (7, 7), strides=(2, 2), name='conv1')(x)
x = BatchNormalization(axis=bn_axis, name='bn_conv1')(x)
x = Activation('relu')(x)
x = MaxPooling2D((3, 3), strides=(2, 2))(x)

x = conv_block(x, 3, [64, 64, 256], stage=2, block='a', strides=(1, 1))
x = identity_block(x, 3, [64, 64, 256], stage=2, block='b')
x = identity_block(x, 3, [64, 64, 256], stage=2, block='c')

x = conv_block(x, 3, [128, 128, 512], stage=3, block='a')
x = identity_block(x, 3, [128, 128, 512], stage=3, block='b')
x = identity_block(x, 3, [128, 128, 512], stage=3, block='c')
x = identity_block(x, 3, [128, 128, 512], stage=3, block='d')

x = conv_block(x, 3, [256, 256, 1024], stage=4, block='a')
x = identity_block(x, 3, [256, 256, 1024], stage=4, block='b')
x = identity_block(x, 3, [256, 256, 1024], stage=4, block='c')
x = identity_block(x, 3, [256, 256, 1024], stage=4, block='d')
x = identity_block(x, 3, [256, 256, 1024], stage=4, block='e')
x = identity_block(x, 3, [256, 256, 1024], stage=4, block='f')

x = conv_block(x, 3, [512, 512, 2048], stage=5, block='a')

```

```
x = identity_block(x, 3, [512, 512, 2048], stage=5, block='b')  
x = identity_block(x, 3, [512, 512, 2048], stage=5, block='c')
```

```
x = AveragePooling2D((7, 7), name='avg_pool')(x)
```

```
if include_top:
```

```
x = Flatten()(x)
```

```
x = Dense(classes, activation='softmax', name='fc1000')(x)
```

```
else:
```

```
if pooling == 'avg':
```

```
x = GlobalAveragePooling2D()(x)
```

```
elif pooling == 'max':
```

```
x = GlobalMaxPooling2D()(x)
```

```
# Ensure that the model takes into account
```

```
# any potential predecessors of `input_tensor`.
```

```
if input_tensor is not None:
```

```
inputs = get_source_inputs(input_tensor)
```

```
else:
```

```
inputs = img_input
```

```
# Create model.
```

```
model = Model(inputs, x, name='resnet50')
```

```
# load weights
```

```
if weights == 'imagenet':
```

```
if include_top:
```

```
weights_path = get_file('resnet50_weights_tf_dim_ordering_tf_kernels.h5',  
                        WEIGHTS_PATH,  
                        cache_subdir='models',  
                        md5_hash='a7b3fe01876f51b976af0dea6bc144eb')
```

```
else:
```

```

    weights_path =
get_file('resnet50_weights_tf_dim_ordering_tf_kernels_notop.h5',
        WEIGHTS_PATH_NO_TOP,
        cache_subdir='models',
        md5_hash='a268eb855778b3df3c7506639542a6af')
model.load_weights(weights_path)
if K.backend() == 'theano':
    layer_utils.convert_all_kernels_in_model(model)

if K.image_data_format() == 'channels_first':
    if include_top:
        maxpool = model.get_layer(name='avg_pool')
        shape = maxpool.output_shape[1:]
        dense = model.get_layer(name='fc1000')
        layer_utils.convert_dense_weights_data_format(dense, shape,
'channels_first')

    if K.backend() == 'tensorflow':
        warnings.warn('You are using the TensorFlow backend, yet you '
            'are using the Theano '
            'image data format convention '
            '('image_data_format="channels_first"). '
            'For best performance, set '
            '`image_data_format="channels_last"` in '
            'your Keras config '
            'at ~/.keras/keras.json.')
    return model

def load_preprocess_training_batch(X_train):

    new = []

```

```
for item in X_train:
    tmpFeature = skimage.transform.resize(item, (224, 224), mode='constant')
    new.append(tmpFeature)
```

```
return new
```

```
def preprocess_data(X_train):
```

```
    for item in X_train:
        item = np.expand_dims(item, axis=0)
        item = preprocess_input(item)
```

```
    return X_train
```

```
# CIFAR-10 DATASET
```

```
(X_train, y_train) , (X_test, y_test) = keras.datasets.cifar10.load_data()
```

```
X_train = X_train[0:2000]
```

```
y_train = y_train[0:2000]
```

```
X_test = X_test[0:2000]
```

```
y_test = y_test[0:2000]
```

```
X_train_resized = load_preprocess_training_batch(X_train)
```

```
X_test_resized = load_preprocess_training_batch(X_test)
```

```
X_train_resized = np.array(X_train_resized)
```

```
X_test_resized = np.array(X_test_resized)
```

```
X_train_resized = X_train_resized / 255
```

```
X_test_resized = X_test_resized / 255
```

```
X_train_resized = preprocess_data(X_train_resized)
```

```
X_test_resized = preprocess_data(X_test_resized)
```

```
model = ResNet50(include_top=True, weights='imagenet')
```

```
model.compile(optimizer='SGD',  
              loss='sparse_categorical_crossentropy',  
              metrics=['accuracy'])
```

```
history = model.fit(X_train_resized, y_train, epochs=5)
```

```
model.evaluate(X_test_resized, y_test)
```

```
# MNIST Dataset
```

```
(X_train, y_train) , (X_test, y_test) = keras.datasets.mnist.load_data()
```

```
X_train = X_train[0:2000]
```

```
y_train = y_train[0:2000]
```

```
X_test = X_test[0:2000]
```

```
y_test = y_test[0:2000]
```

```
X_train_resized = load_preprocess_training_batch(X_train)
```

```
X_test_resized = load_preprocess_training_batch(X_test)
```

```
X_train_resized = np.array(X_train_resized)
```

```
X_test_resized = np.array(X_test_resized)
```

```
X_train_resized = X_train_resized / 255.0
```

```
X_test_resized = X_test_resized / 255.0
```

```
X_train_resized = preprocess_data(X_train_resized)
```

```
X_test_resized = preprocess_data(X_test_resized)

import cv2

X_train_new = list()

for i in range(len(X_train_resized)):
    g = X_train_resized[i]
    X_train_new.append(cv2.merge([g,g,g]))

X_train_new = np.asarray(X_train_new,dtype=np.float32)

X_test_new = list()

for i in range(len(X_test_resized)):
    g = X_test_resized[i]
    X_test_new.append(cv2.merge([g,g,g]))

X_test_new = np.asarray(X_test_new,dtype=np.float32)

model = ResNet50(include_top=True, weights='imagenet')

model.compile(optimizer='SGD',
               loss='sparse_categorical_crossentropy',
               metrics=['accuracy'])

history = model.fit(X_train_new, y_train, epochs=10)

model.evaluate(X_test_new, y_test)

# SAVEE Dataset
```

```
!unzip "/content/drive/MyDrive/SaveeDataset.zip"
```

```
import librosa
```

```
import numpy as np
```

```
input_length = 16000*5
```

```
batch_size = 32
```

```
n_mels = 320
```

```
def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,  
#log_specgram  
    step_size=10, eps=1e-10):
```

```
    mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,  
n_mels= n_mels)  
    mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40  
  
    return mel_db.T
```

```
def load_audio_file(file_path, input_length=input_length):  
    data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000  
    if len(data)>input_length:  
        max_offset = len(data)-input_length  
  
        offset = np.random.randint(max_offset)  
  
        data = data[offset:(input_length+offset)]  
  
    else:
```

```
if input_length > len(data):
    max_offset = input_length - len(data)

    offset = np.random.randint(max_offset)
else:
    offset = 0
data = np.pad(data, (offset, input_length - len(data) - offset), "constant")
```

```
data = preprocess_audio_mel_T(data)
return data
```

Preprocessing the dataset

```
import os
from scipy.io import wavfile
import librosa
import matplotlib.pyplot as plt
import numpy as np
import cv2
```

```
rootDirectory = "/content/AudioData/"
personNames = ["DC","JE","JK","KL"]
```

```
classes = ["a" , "d" , "f" , "h" , "n" , "sa" , "su" ]
```

```
X = list()
y = list()
```

```
for person in personNames:
    directory = os.path.join(rootDirectory,person)
    for filename in os.listdir(directory):
        filePath = os.path.join(directory, filename)
        a = load_audio_file(file_path=filePath)
```



```
data = cv2.merge([a,a,a])  
# data = np.reshape(data, data.shape + (1,))  
if(filename[0:1] in classes):  
X.append(data)  
y.append(classes.index(filename[0:1]))  
elif(filename[0:2] in classes):  
X.append(data)  
y.append(classes.index(filename[0:2]))
```

```
X = np.asarray(X, dtype=np.float32)  
y = np.asarray(y, dtype=np.float32)
```

```
import pandas as pd  
import numpy as np  
import matplotlib.pyplot as plt  
import tensorflow as tf
```

```
# dataset preparation
```

```
from tensorflow.keras import datasets, layers, models  
from sklearn.model_selection import train_test_split  
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.5, train_size= 0.5  
,random_state=10)
```

```
X_train_resized = load_preprocess_training_batch(X_train)  
X_test_resized = load_preprocess_training_batch(X_test)
```

```
X_train_resized = np.array(X_train_resized)  
X_test_resized = np.array(X_test_resized)
```

```
X_train_resized = preprocess_data(X_train_resized)  
X_test_resized = preprocess_data(X_test_resized)
```

```
model = ResNet50(include_top=True, weights='imagenet')

model.compile(optimizer='SGD',
              loss='sparse_categorical_crossentropy',
              metrics=['accuracy'])

history = model.fit(X_train_resized, y_train, epochs=10)

model.evaluate(X_test_resized, y_test)

# EmoDB Dataset

!unzip "/content/drive/MyDrive/EmoDB.zip"

import librosa
import numpy as np

input_length = 16000*5

batch_size = 32

n_mels = 320

def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,
#log_specgram
    step_size=10, eps=1e-10):

    mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,
n_mels= n_mels)
    mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40
```

```
return mel_db.T
```

```
def load_audio_file(file_path, input_length=input_length):  
    data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000  
    if len(data)>input_length:  
        max_offset = len(data)-input_length  
  
        offset = np.random.randint(max_offset)  
  
        data = data[offset:(input_length+offset)]  
  
    else:  
        if input_length > len(data):  
            max_offset = input_length - len(data)  
  
            offset = np.random.randint(max_offset)  
        else:  
            offset = 0  
        data = np.pad(data, (offset, input_length - len(data) - offset), "constant")  
  
    data = preprocess_audio_mel_T(data)  
    return data
```

```
# Preprocessing the dataset
```

```
import os  
from scipy.io import wavfile  
import librosa  
import matplotlib.pyplot as plt  
import numpy as np  
import cv2
```

```
directory = "/content/wav/"
```

```
classes = ["W" ,"L" ,"E" ,"A" ,"F" ,"T" ,"N" ]
```

```
X = list()
```

```
y = list()
```

```
for filename in os.listdir(directory):
```

```
    filePath = os.path.join(directory, filename)
```

```
    a = load_audio_file(file_path=filePath)
```

```
    data = cv2.merge([a,a,a])
```

```
    if(filename[5:6] in classes):
```

```
        X.append(data)
```

```
        y.append(classes.index(filename[5:6]))
```

```
X = np.asarray(X, dtype=np.float32)
```

```
y = np.asarray(y, dtype=np.float32)
```

```
import pandas as pd
```

```
import numpy as np
```

```
import matplotlib.pyplot as plt
```

```
import tensorflow as tf
```

```
# dataset preparation
```

```
from tensorflow.keras import datasets, layers, models
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.5, train_size= 0.5  
, random_state=10)
```

```
X_train_resized = load_preprocess_training_batch(X_train)
```

```
X_test_resized = load_preprocess_training_batch(X_test)
```

```
X_train_resized = np.array(X_train_resized)
```

```
X_test_resized = np.array(X_test_resized)
```

```
X_train_resized = preprocess_data(X_train_resized)
```

```
X_test_resized = preprocess_data(X_test_resized)
```

```
model = ResNet50(include_top=True, weights='imagenet')
```

```
model.compile(optimizer='SGD',  
              loss='sparse_categorical_crossentropy',  
              metrics=['accuracy'])
```

```
history = model.fit(X_train_resized, y_train, epochs=10)
```

```
model.evaluate(X_test_resized, y_test)#@title
```

```
from google.colab import drive
```

```
drive.mount('/content/drive')
```

```
# CIFAR 10 DATASET
```

```
#@title
```

```
import os
```

```
import tensorflow as tf
```

```
import keras
```

```
from tensorflow.keras import layers
```

```
from tensorflow.keras import Model
```

```
from os import getcwd
```

```
#@title
```

```
cifar10 = tf.keras.datasets.cifar10
```

```
(training_images, training_labels), (test_images, test_labels) = cifar10.load_data()
```

#@title

print(len(training_images))

print(len(test_images))

#@title

training_images = training_images.reshape(50000, 1024, 3)

training_images = training_images[0:10000]

training_labels = training_labels[0:10000]

training_images = training_images/255.0

test_images = test_images.reshape(10000, 1024, 3)

test_images = test_images[0:5000]

test_labels = test_labels[0:5000]

test_images = test_images/255.0

#@title

model = tf.keras.models.Sequential([

tf.keras.layers.Bidirectional(tf.keras.layers.LSTM(32, input_shape=(1024,3),
return_sequences=True)),

tf.keras.layers.Bidirectional(tf.keras.layers.LSTM(32)),

tf.keras.layers.Flatten(),

tf.keras.layers.Dense(64, activation='relu'),

tf.keras.layers.Dense(10, activation='softmax')

])

#@title

model.compile(optimizer='adam',

loss='sparse_categorical_crossentropy',

metrics=['accuracy'])

history = model.fit(training_images, training_labels, batch_size = 50, epochs=10)

#@title

model.evaluate(test_images, test_labels)

MNIST DATASET

#@title

import torch

#@title

Device configuration

device = torch.device('cuda' if torch.cuda.is_available() else 'cpu')

device

#@title

from torchvision import datasets

from torchvision.transforms import ToTensor

train_data = datasets.MNIST(

root = 'data',

train = True,

transform = ToTensor(),

download = True,

)

test_data = datasets.MNIST(

root = 'data',

train = False,

transform = ToTensor()

)

#@title

print(train_data)

```
#@title  
print(test_data)
```

```
#@title  
print(train_data.data.size())
```

```
#@title  
print(train_data.targets.size())
```

```
#@title  
print(train_data.data[0])
```

```
#@title  
import matplotlib.pyplot as plt  
plt.imshow(train_data.data[0], cmap='gray')  
plt.title('%i' % train_data.targets[0])  
plt.show()
```

```
#@title  
figure = plt.figure(figsize=(10, 8))  
cols, rows = 5, 5  
for i in range(1, cols * rows + 1):  
    sample_idx = torch.randint(len(train_data), size=(1,)).item()  
    img, label = train_data[sample_idx]  
    figure.add_subplot(rows, cols, i)  
    plt.title(label)
```



```
plt.axis("off")
plt.imshow(img.squeeze(), cmap="gray")
plt.show()
```

```
##@title
```

```
from torch.utils.data import DataLoader
```

```
loaders = {
    'train' : torch.utils.data.DataLoader(train_data,
                                           batch_size=100,
                                           shuffle=True,
                                           num_workers=1),

    'test' : torch.utils.data.DataLoader(test_data,
                                           batch_size=100,
                                           shuffle=True,
                                           num_workers=1),
}
loaders
```

```
##@title
```

```
from torch import nn
```

```
import torch.nn.functional as F
```

```
##@title
```

```
sequence_length = 28
```

```
input_size = 28
```

```
hidden_size = 128
```

```
num_layers = 2
```

```
num_classes = 10
```

```
batch_size = 100
```

```
num_epochs = 2  
learning_rate = 0.01
```

```
#@title  
class RNN(nn.Module):  
    pass  
model = RNN().to(device)  
print(model)
```

```
#@title  
class RNN(nn.Module):  
  
    def __init__(self, input_size, hidden_size, num_layers, num_classes):  
        super(RNN, self).__init__()  
        self.hidden_size = hidden_size  
        self.num_layers = num_layers  
        self.lstm = nn.LSTM(input_size, hidden_size, num_layers, batch_first=True)  
        self.fc = nn.Linear(hidden_size, num_classes)  
        pass  
  
    def forward(self, x):  
        # Set initial hidden and cell states  
        h0 = torch.zeros(self.num_layers, x.size(0), self.hidden_size).to(device)  
        c0 = torch.zeros(self.num_layers, x.size(0), self.hidden_size).to(device)  
        # Passing in the input and hidden state into the model and obtaining outputs  
        out, hidden = self.lstm(x, (h0, c0)) # out: tensor of shape (batch_size,  
seq_length, hidden_size)  
  
        #Reshaping the outputs such that it can be fit into the fully connected layer  
        out = self.fc(out[:, -1, :])
```

return out

pass

pass

model = RNN(input_size, hidden_size, num_layers, num_classes).to(device)

print(model)

#@title

loss_func = nn.CrossEntropyLoss()

loss_func

#@title

from torch import optim

optimizer = optim.Adam(model.parameters(), lr = 0.01)

optimizer

#@title

def train(num_epochs, model, loaders):

Train the model

total_step = len(loaders['train'])

for epoch in range(num_epochs):

for i, (images, labels) in enumerate(loaders['train']):

images = images.reshape(-1, sequence_length, input_size).to(device)

labels = labels.to(device)

Forward pass

outputs = model(images)

loss = loss_func(outputs, labels)

```

# Backward and optimize
optimizer.zero_grad()
loss.backward()
optimizer.step()

if (i+1) % 100 == 0:
    print ('Epoch [{}/{}], Step [{}/{}], Loss: {:.4f}'
        .format(epoch + 1, num_epochs, i + 1, total_step, loss.item()))
    pass

pass
pass
train(num_epochs, model, loaders)

# @title
# Test the model
model.eval()
with torch.no_grad():
    correct = 0
    total = 0
    for images, labels in loaders['test']:
        images = images.reshape(-1, sequence_length, input_size).to(device)
        labels = labels.to(device)
        outputs = model(images)
        _, predicted = torch.max(outputs.data, 1)
        total = total + labels.size(0)
        correct = correct + (predicted == labels).sum().item()
print('Test Accuracy of the model on the 10000 test images: {} %'.format(100 * correct / total))

# SAVEE Dataset

```

```

#@title
!unzip "/content/drive/MyDrive/SaveeDataset.zip"

#@title
import librosa
import numpy as np

input_length = 16000*5

batch_size = 32

n_mels = 320

def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,
#log_specgram
    step_size=10, eps=1e-10):

    mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,
n_mels= n_mels)
    mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40

    return mel_db.T

def load_audio_file(file_path, input_length=input_length):
    data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000
    if len(data)>input_length:
        max_offset = len(data)-input_length

        offset = np.random.randint(max_offset)

        data = data[offset:(input_length+offset)]

```

```
else:
    if input_length > len(data):
        max_offset = input_length - len(data)

        offset = np.random.randint(max_offset)
    else:
        offset = 0
    data = np.pad(data, (offset, input_length - len(data) - offset), "constant")

data = preprocess_audio_mel_T(data)
return data
```

```
# @title
# Preprocessing the dataset
import os
from scipy.io import wavfile
import librosa
import matplotlib.pyplot as plt
import numpy as np
import cv2

rootDirectory = "/content/AudioData/"
personNames = ["DC", "JE", "JK", "KL"]

classes = ["a", "d", "f", "h", "n", "sa", "su"]

X = list()
y = list()

for person in personNames:
    directory = os.path.join(rootDirectory, person)
```

```
for filename in os.listdir(directory):  
    filePath = os.path.join(directory, filename)  
    data = load_audio_file(file_path=filePath)  
    # data = cv2.merge([a,a,a])  
    if(filename[0:1] in classes):  
        X.append(data)  
        y.append(classes.index(filename[0:1]))  
    elif(filename[0:2] in classes):  
        X.append(data)  
        y.append(classes.index(filename[0:2]))
```

#@title

```
X = np.asarray(X, dtype=np.float32)  
y = np.asarray(y, dtype=np.float32)
```

#@title

```
X.shape , y.shape
```

#@title

```
import pandas as pd  
import numpy as np  
import matplotlib.pyplot as plt  
import tensorflow as tf
```

dataset preparation

```
from tensorflow.keras import datasets, layers, models  
from sklearn.model_selection import train_test_split  
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.4, train_size= 0.6  
, random_state=10)
```

#@title

```

import os
import tensorflow as tf
import keras
from tensorflow.keras import layers
from tensorflow.keras import Model
from os import getcwd

#@title
model = tf.keras.models.Sequential([
    tf.keras.layers.Bidirectional(tf.keras.layers.LSTM(32, input_shape=(157,320),
return_sequences=True)),
    tf.keras.layers.Bidirectional(tf.keras.layers.LSTM(32)),
    tf.keras.layers.Flatten(),
    tf.keras.layers.Dense(64, activation='relu'),
    tf.keras.layers.Dense(10, activation='softmax')
])

#@title
model.compile(optimizer='adam',
              loss='sparse_categorical_crossentropy',
              metrics=['accuracy'])

history = model.fit(X_train,y_train, batch_size = 50, epochs=50)

#@title
model.evaluate(X_test, y_test)

# EmoDB Dataset

#@title
!unzip "/content/drive/MyDrive/EmoDB.zip"

```



```
#@title
import librosa
import numpy as np

input_length = 16000*5

batch_size = 32

n_mels = 320

def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,
#log_specgram
    step_size=10, eps=1e-10):

    mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,
n_mels= n_mels)
    mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40

    return mel_db.T

def load_audio_file(file_path, input_length=input_length):
    data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000
    if len(data)>input_length:
        max_offset = len(data)-input_length

        offset = np.random.randint(max_offset)

        data = data[offset:(input_length+offset)]

    else:
        if input_length > len(data):
```

```
max_offset = input_length - len(data)
```

```
offset = np.random.randint(max_offset)
```

```
else:
```

```
offset = 0
```

```
data = np.pad(data, (offset, input_length - len(data) - offset), "constant")
```

```
data = preprocess_audio_mel_T(data)
```

```
return data
```

```
#@title
```

```
# Preprocessing the dataset
```

```
import os
```

```
from scipy.io import wavfile
```

```
import librosa
```

```
import matplotlib.pyplot as plt
```

```
import numpy as np
```

```
import cv2
```

```
directory = "/content/wav/"
```

```
classes = ["W" , "L" , "E" , "A" , "F" , "T" , "N" ]
```

```
X = list()
```

```
y = list()
```

```
for filename in os.listdir(directory):
```

```
    filePath = os.path.join(directory, filename)
```

```
    data = load_audio_file(file_path=filePath)
```

```
    if(filename[5:6] in classes):
```

```
        X.append(data)
```

```
        y.append(classes.index(filename[5:6]))
```

#@title

X = np.asarray(X, dtype=np.float32)

y = np.asarray(y, dtype=np.float32)

#@title

import pandas as pd

import numpy as np

import matplotlib.pyplot as plt

import tensorflow as tf

dataset preparation

from tensorflow.keras import datasets, layers, models

from sklearn.model_selection import train_test_split

**X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, train_size= 0.7
, random_state=10)**

#@title

model = tf.keras.models.Sequential([

**tf.keras.layers.Bidirectional(tf.keras.layers.LSTM(32, input_shape=(157,320),
return_sequences=True)),**

tf.keras.layers.Bidirectional(tf.keras.layers.LSTM(32)),

tf.keras.layers.Flatten(),

tf.keras.layers.Dense(64, activation='relu'),

tf.keras.layers.Dense(10, activation='softmax')

])

#@title

model.compile(optimizer='adam',

loss='sparse_categorical_crossentropy',

metrics=['accuracy'])

```
history = model.fit(X_train,y_train, batch_size = 50, epochs=50)
```

```
# @title
```

```
model.evaluate(X_test, y_test)from google.colab import drive  
drive.mount('/content/drive')
```

```
# CIFAR 10 DATASET
```

```
def inception_module(x,  
    filters_1x1,  
    filters_3x3_reduce,  
    filters_3x3,  
    filters_5x5_reduce,  
    filters_5x5,  
    filters_pool_proj,  
    name=None):
```

```
    conv_1x1 = Conv2D(filters_1x1, (1, 1), padding='same', activation='relu',  
kernel_initializer=kernel_init, bias_initializer=bias_init)(x)
```

```
    conv_3x3_reduce = Conv2D(filters_3x3_reduce, (1, 1), padding='same',  
activation='relu', kernel_initializer=kernel_init, bias_initializer=bias_init)(x)  
    conv_3x3 = Conv2D(filters_3x3, (3, 3), padding='same', activation='relu',  
kernel_initializer=kernel_init, bias_initializer=bias_init)(conv_3x3_reduce)
```

```
    conv_5x5_reduce = Conv2D(filters_5x5_reduce, (1, 1), padding='same',  
activation='relu', kernel_initializer=kernel_init, bias_initializer=bias_init)(x)  
    conv_5x5 = Conv2D(filters_5x5, (5, 5), padding='same', activation='relu',  
kernel_initializer=kernel_init, bias_initializer=bias_init)(conv_5x5_reduce)
```

```
    pool_proj = MaxPool2D((3, 3), strides=(1, 1), padding='same')(x)
```

```
pool_proj = Conv2D(filters_pool_proj, (1, 1), padding='same',  
activation='relu', kernel_initializer=kernel_init,  
bias_initializer=bias_init)(pool_proj)
```

```
output = concatenate([conv_1x1, conv_3x3, conv_5x5, pool_proj], axis=3,  
name=name)
```

```
return output
```

```
kernel_init = keras.initializers.glorot_uniform()  
bias_init = keras.initializers.Constant(value=0.2)
```

```
input_layer = Input(shape=(224, 224, 3))
```

```
x = Conv2D(64, (7, 7), padding='same', strides=(2, 2), activation='relu',  
name='conv_1_7x7/2', kernel_initializer=kernel_init,  
bias_initializer=bias_init)(input_layer)
```

```
x = MaxPool2D((3, 3), padding='same', strides=(2, 2),  
name='max_pool_1_3x3/2')(x)
```

```
x = Conv2D(64, (1, 1), padding='same', strides=(1, 1), activation='relu',  
name='conv_2a_3x3/1')(x)
```

```
x = Conv2D(192, (3, 3), padding='same', strides=(1, 1), activation='relu',  
name='conv_2b_3x3/1')(x)
```

```
x = MaxPool2D((3, 3), padding='same', strides=(2, 2),  
name='max_pool_2_3x3/2')(x)
```

```
x = inception_module(x,  
    filters_1x1=64,  
    filters_3x3_reduce=96,  
    filters_3x3=128,  
    filters_5x5_reduce=16,  
    filters_5x5=32,
```

```
filters_pool_proj=32,  
name='inception_3a')
```

```
x = inception_module(x,  
    filters_1x1=128,  
    filters_3x3_reduce=128,  
    filters_3x3=192,  
    filters_5x5_reduce=32,  
    filters_5x5=96,  
    filters_pool_proj=64,  
    name='inception_3b')
```

```
x = MaxPool2D((3, 3), padding='same', strides=(2, 2),  
name='max_pool_3_3x3/2')(x)
```

```
x = inception_module(x,  
    filters_1x1=192,  
    filters_3x3_reduce=96,  
    filters_3x3=208,  
    filters_5x5_reduce=16,  
    filters_5x5=48,  
    filters_pool_proj=64,  
    name='inception_4a')
```

```
x1 = AveragePooling2D((5, 5), strides=3)(x)  
x1 = Conv2D(128, (1, 1), padding='same', activation='relu')(x1)  
x1 = Flatten()(x1)  
x1 = Dense(1024, activation='relu')(x1)  
x1 = Dropout(0.7)(x1)  
x1 = Dense(10, activation='softmax', name='auxilliary_output_1')(x1)
```

```
x = inception_module(x,  
    filters_1x1=160,  
    filters_3x3_reduce=112,  
    filters_3x3=224,  
    filters_5x5_reduce=24,  
    filters_5x5=64,  
    filters_pool_proj=64,  
    name='inception_4b')
```

```
x = inception_module(x,  
    filters_1x1=128,  
    filters_3x3_reduce=128,  
    filters_3x3=256,  
    filters_5x5_reduce=24,  
    filters_5x5=64,  
    filters_pool_proj=64,  
    name='inception_4c')
```

```
x = inception_module(x,  
    filters_1x1=112,  
    filters_3x3_reduce=144,  
    filters_3x3=288,  
    filters_5x5_reduce=32,  
    filters_5x5=64,  
    filters_pool_proj=64,  
    name='inception_4d')
```

```
x2 = AveragePooling2D((5, 5), strides=3)(x)  
x2 = Conv2D(128, (1, 1), padding='same', activation='relu')(x2)  
x2 = Flatten()(x2)  
x2 = Dense(1024, activation='relu')(x2)
```

```
x2 = Dropout(0.7)(x2)
```

```
x2 = Dense(10, activation='softmax', name='auxilliary_output_2')(x2)
```

```
x = inception_module(x,  
    filters_1x1=256,  
    filters_3x3_reduce=160,  
    filters_3x3=320,  
    filters_5x5_reduce=32,  
    filters_5x5=128,  
    filters_pool_proj=128,  
    name='inception_4e')
```

```
x = MaxPool2D((3, 3), padding='same', strides=(2, 2),  
name='max_pool_4_3x3/2')(x)
```

```
x = inception_module(x,  
    filters_1x1=256,  
    filters_3x3_reduce=160,  
    filters_3x3=320,  
    filters_5x5_reduce=32,  
    filters_5x5=128,  
    filters_pool_proj=128,  
    name='inception_5a')
```

```
x = inception_module(x,  
    filters_1x1=384,  
    filters_3x3_reduce=192,  
    filters_3x3=384,  
    filters_5x5_reduce=48,  
    filters_5x5=128,  
    filters_pool_proj=128,  
    name='inception_5b')
```



```
x = GlobalAveragePooling2D(name='avg_pool_5_3x3/1')(x)
```

```
x = Dropout(0.4)(x)
```

```
x = Dense(10, activation='softmax', name='output')(x)
```

```
import keras
```

```
from keras.layers.core import Layer
```

```
import keras.backend as K
```

```
import tensorflow as tf
```

```
from keras.datasets import cifar10
```

```
from keras.models import Model
```

```
from keras.layers import Conv2D, MaxPool2D, \  
    Dropout, Dense, Input, concatenate, \  
    GlobalAveragePooling2D, AveragePooling2D, \  
    Flatten
```

```
import cv2
```

```
import numpy as np
```

```
from keras.datasets import cifar10
```

```
from keras import backend as K
```

```
from keras.utils import np_utils
```

```
import math
```

```
from tensorflow.keras.optimizers import SGD
```

```
from keras.callbacks import LearningRateScheduler
```

```
num_classes = 10
```

```

def load_cifar10_data(img_rows, img_cols):

    # Load cifar10 training and validation sets
    (X_train, Y_train), (X_valid, Y_valid) = cifar10.load_data()

    X_train = X_train[0:5000]
    Y_train = Y_train[0:5000]
    X_valid = X_valid[0:2000]
    Y_valid = Y_valid[0:2000]

    # Resize training images
    X_train = np.array([cv2.resize(img, (img_rows, img_cols)) for img in
X_train[:, :, :, :]])
    X_valid = np.array([cv2.resize(img, (img_rows, img_cols)) for img in
X_valid[:, :, :, :]])

    # Transform targets to keras compatible format
    Y_train = np_utils.to_categorical(Y_train, num_classes)
    Y_valid = np_utils.to_categorical(Y_valid, num_classes)

    X_train = X_train.astype('float32')
    X_valid = X_valid.astype('float32')

    # preprocess data
    X_train = X_train / 255.0
    X_valid = X_valid / 255.0

    return X_train, Y_train, X_valid, Y_valid

X_train, y_train, X_test, y_test = load_cifar10_data(224, 224)

```

```
model = Model(input_layer, [x, x1, x2], name='inception_v1')
```

```
model.summary()
```

```
epochs = 10
```

```
initial_lrate = 0.01
```

```
def decay(epoch, steps=100):
```

```
    initial_lrate = 0.01
```

```
    drop = 0.96
```

```
    epochs_drop = 8
```

```
    lrate = initial_lrate * math.pow(drop, math.floor((1+epoch)/epochs_drop))
```

```
    return lrate
```

```
sgd = SGD(learning_rate=initial_lrate, momentum=0.9, nesterov=False)
```

```
lr_sc = LearningRateScheduler(decay, verbose=1)
```

```
model.compile(loss=['categorical_crossentropy', 'categorical_crossentropy',  
'categorical_crossentropy'], loss_weights=[1, 0.3, 0.3], optimizer=sgd,  
metrics=['accuracy'])
```

```
history = model.fit(X_train, [y_train, y_train, y_train], validation_data=(X_test,  
[y_test, y_test, y_test]), epochs=epochs, batch_size=256, callbacks=[lr_sc])
```

```
# MNIST DATASET
```

```
import tensorflow as tf
```

```
import matplotlib.pyplot as plt
```

```
from tensorflow.keras import datasets, layers, models, losses, Model
```

```
(x_train, y_train), (x_test, y_test)=tf.keras.datasets.mnist.load_data()  
x_train = tf.pad(x_train, [[0, 0], [2,2], [2,2]])/255  
x_test = tf.pad(x_test, [[0, 0], [2,2], [2,2]])/255  
x_train = tf.expand_dims(x_train, axis=3, name=None)  
x_test = tf.expand_dims(x_test, axis=3, name=None)  
x_train = tf.repeat(x_train, 3, axis=3)  
x_test = tf.repeat(x_test, 3, axis=3)  
x_val = x_train[-2000::, :, :]  
y_val = y_train[-2000:]  
x_train = x_train[:-2000, :, :]  
y_train = y_train[:-2000]
```

```
def inception(x,  
             filters_1x1,  
             filters_3x3_reduce,  
             filters_3x3,  
             filters_5x5_reduce,  
             filters_5x5,  
             filters_pool):  
    path1 = layers.Conv2D(filters_1x1, (1, 1), padding='same', activation='relu')(x)  
  
    path2 = layers.Conv2D(filters_3x3_reduce, (1, 1), padding='same',  
activation='relu')(x)  
    path2 = layers.Conv2D(filters_3x3, (1, 1), padding='same',  
activation='relu')(path2)  
  
    path3 = layers.Conv2D(filters_5x5_reduce, (1, 1), padding='same',  
activation='relu')(x)  
    path3 = layers.Conv2D(filters_5x5, (1, 1), padding='same',  
activation='relu')(path3)
```

```
path4 = layers.MaxPool2D((3, 3), strides=(1, 1), padding='same')(x)
path4 = layers.Conv2D(filters_pool, (1, 1), padding='same',
activation='relu')(path4)
```

```
return tf.concat([path1, path2, path3, path4], axis=3)
```

```
inp = layers.Input(shape=(32, 32, 3))
input_tensor = layers.experimental.preprocessing.Resizing(224, 224,
interpolation="bilinear", input_shape=x_train.shape[1:])(inp)
```

```
x = layers.Conv2D(64, 7, strides=2, padding='same',
activation='relu')(input_tensor)
x = layers.MaxPooling2D(3, strides=2)(x)
```

```
x = layers.Conv2D(64, 1, strides=1, padding='same', activation='relu')(x)
x = layers.Conv2D(192, 3, strides=1, padding='same', activation='relu')(x)
```

```
x = layers.MaxPooling2D(3, strides=2)(x)
```

```
x = inception(x,
    filters_1x1=64,
    filters_3x3_reduce=96,
    filters_3x3=128,
    filters_5x5_reduce=16,
    filters_5x5=32,
    filters_pool=32)
```

```
x = inception(x,
    filters_1x1=128,
    filters_3x3_reduce=128,
```

```
filters_3x3=192,  
filters_5x5_reduce=32,  
filters_5x5=96,  
filters_pool=64)
```

```
x = layers.MaxPooling2D(3, strides=2)(x)
```

```
x = inception(x,  
filters_1x1=192,  
filters_3x3_reduce=96,  
filters_3x3=208,  
filters_5x5_reduce=16,  
filters_5x5=48,  
filters_pool=64)
```

```
aux1 = layers.AveragePooling2D((5, 5), strides=3)(x)  
aux1 = layers.Conv2D(128, 1, padding='same', activation='relu')(aux1)  
aux1 = layers.Flatten()(aux1)  
aux1 = layers.Dense(1024, activation='relu')(aux1)  
aux1 = layers.Dropout(0.7)(aux1)  
aux1 = layers.Dense(10, activation='softmax')(aux1)
```

```
x = inception(x,  
filters_1x1=160,  
filters_3x3_reduce=112,  
filters_3x3=224,  
filters_5x5_reduce=24,  
filters_5x5=64,  
filters_pool=64)
```

```
x = inception(x,  
filters_1x1=128,
```

```
filters_3x3_reduce=128,  
filters_3x3=256,  
filters_5x5_reduce=24,  
filters_5x5=64,  
filters_pool=64)
```

```
x = inception(x,  
              filters_1x1=112,  
              filters_3x3_reduce=144,  
              filters_3x3=288,  
              filters_5x5_reduce=32,  
              filters_5x5=64,  
              filters_pool=64)
```

```
aux2 = layers.AveragePooling2D((5, 5), strides=3)(x)  
aux2 = layers.Conv2D(128, 1, padding='same', activation='relu')(aux2)  
aux2 = layers.Flatten()(aux2)  
aux2 = layers.Dense(1024, activation='relu')(aux2)  
aux2 = layers.Dropout(0.7)(aux2)  
aux2 = layers.Dense(10, activation='softmax')(aux2)
```

```
x = inception(x,  
              filters_1x1=256,  
              filters_3x3_reduce=160,  
              filters_3x3=320,  
              filters_5x5_reduce=32,  
              filters_5x5=128,  
              filters_pool=128)
```

```
x = layers.MaxPooling2D(3, strides=2)(x)
```

```
x = inception(x,
```

```
filters_1x1=256,  
filters_3x3_reduce=160,  
filters_3x3=320,  
filters_5x5_reduce=32,  
filters_5x5=128,  
filters_pool=128)
```

```
x = inception(x,  
filters_1x1=384,  
filters_3x3_reduce=192,  
filters_3x3=384,  
filters_5x5_reduce=48,  
filters_5x5=128,  
filters_pool=128)
```

```
x = layers.GlobalAveragePooling2D()(x)
```

```
x = layers.Dropout(0.4)(x)  
out = layers.Dense(10, activation='softmax')(x)
```

```
model = Model(inputs = inp, outputs = [out, aux1, aux2])
```

```
model.compile(optimizer='adam', loss=[losses.sparse_categorical_crossentropy,  
losses.sparse_categorical_crossentropy, losses.sparse_categorical_crossentropy],  
loss_weights=[1, 0.3, 0.3], metrics=['accuracy'])
```

```
history = model.fit(x_train, [y_train, y_train, y_train], validation_data=(x_val,  
[y_val, y_val, y_val]), batch_size=64, epochs=10)
```

```
fig, axs = plt.subplots(2, 1, figsize=(15,15))
```



```
axs[0].plot(history.history['loss'])
axs[0].plot(history.history['val_loss'])
axs[0].title.set_text('Training Loss vs Validation Loss')
axs[0].set_xlabel('Epochs')
axs[0].set_ylabel('Loss')
axs[0].legend(['Train', 'Val'])
```

```
model.evaluate(x_test, y_test)
```

```
# SAVEE Dataset
```

```
!unzip "/content/drive/MyDrive/SaveeDataset.zip"
```

```
import librosa
import numpy as np
```

```
input_length = 16000*5
```

```
batch_size = 32
```

```
n_mels = 320
```

```
def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,
#log_specgram
    step_size=10, eps=1e-10):
```

```
    mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,
n_mels= n_mels)
    mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40
```

```
return mel_db.T
```

```
def load_audio_file(file_path, input_length=input_length):  
    data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000  
    if len(data)>input_length:  
        max_offset = len(data)-input_length  
  
        offset = np.random.randint(max_offset)  
  
        data = data[offset:(input_length+offset)]  
  
    else:  
        if input_length > len(data):  
            max_offset = input_length - len(data)  
  
            offset = np.random.randint(max_offset)  
        else:  
            offset = 0  
        data = np.pad(data, (offset, input_length - len(data) - offset), "constant")  
  
    data = preprocess_audio_mel_T(data)  
    return data
```

```
# Preprocessing the dataset
```

```
import os
```

```
from scipy.io import wavfile
```

```
import librosa
```

```
import matplotlib.pyplot as plt
```

```
import numpy as np
```

```
import cv2
```

```
rootDirectory = "/content/AudioData/"  
personNames = ["DC","JE","JK","KL"]  
  
classes = ["a" , "d" , "f" , "h" , "n" , "sa" , "su" ]
```

```
X = list()  
y = list()
```

```
for person in personNames:  
    directory = os.path.join(rootDirectory,person)  
    for filename in os.listdir(directory):  
        filePath = os.path.join(directory, filename)  
        a = load_audio_file(file_path=filePath)  
        data = cv2.merge([a,a,a])  
        if(filename[0:1] in classes):  
            X.append(data)  
            y.append(classes.index(filename[0:1]))  
        elif(filename[0:2] in classes):  
            X.append(data)  
            y.append(classes.index(filename[0:2]))
```

```
X = np.asarray(X, dtype=np.float32)  
y = np.asarray(y, dtype=np.float32)
```

```
import pandas as pd  
import numpy as np  
import matplotlib.pyplot as plt  
import tensorflow as tf
```

```
# dataset preparation
```

```
from tensorflow.keras import datasets, layers, models
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.4, train_size= 0.6
,random_state=10)
```

```
import tensorflow as tf
import matplotlib.pyplot as plt
from tensorflow.keras import datasets, layers, models, losses, Model
```

```
def inception(x,
              filters_1x1,
              filters_3x3_reduce,
              filters_3x3,
              filters_5x5_reduce,
              filters_5x5,
              filters_pool):
    path1 = layers.Conv2D(filters_1x1, (1, 1), padding='same', activation='relu')(x)

    path2 = layers.Conv2D(filters_3x3_reduce, (1, 1), padding='same',
activation='relu')(x)
    path2 = layers.Conv2D(filters_3x3, (1, 1), padding='same',
activation='relu')(path2)

    path3 = layers.Conv2D(filters_5x5_reduce, (1, 1), padding='same',
activation='relu')(x)
    path3 = layers.Conv2D(filters_5x5, (1, 1), padding='same',
activation='relu')(path3)

    path4 = layers.MaxPool2D((3, 3), strides=(1, 1), padding='same')(x)
    path4 = layers.Conv2D(filters_pool, (1, 1), padding='same',
activation='relu')(path4)
```

```
return tf.concat([path1, path2, path3, path4], axis=3)
```

```
inp = layers.Input(shape=(157, 320, 3))
```

```
input_tensor = layers.experimental.preprocessing.Resizing(224, 224,  
interpolation='bilinear', input_shape=X_train.shape[1:])(inp)
```

```
x = layers.Conv2D(64, 7, strides=2, padding='same',  
activation='relu')(input_tensor)
```

```
x = layers.MaxPooling2D(3, strides=2)(x)
```

```
x = layers.Conv2D(64, 1, strides=1, padding='same', activation='relu')(x)
```

```
x = layers.Conv2D(192, 3, strides=1, padding='same', activation='relu')(x)
```

```
x = layers.MaxPooling2D(3, strides=2)(x)
```

```
x = inception(x,  
             filters_1x1=64,  
             filters_3x3_reduce=96,  
             filters_3x3=128,  
             filters_5x5_reduce=16,  
             filters_5x5=32,  
             filters_pool=32)
```

```
x = inception(x,  
             filters_1x1=128,  
             filters_3x3_reduce=128,  
             filters_3x3=192,  
             filters_5x5_reduce=32,  
             filters_5x5=96,  
             filters_pool=64)
```

```
x = layers.MaxPooling2D(3, strides=2)(x)
```

```
x = inception(x,  
    filters_1x1=192,  
    filters_3x3_reduce=96,  
    filters_3x3=208,  
    filters_5x5_reduce=16,  
    filters_5x5=48,  
    filters_pool=64)
```

```
aux1 = layers.AveragePooling2D((5, 5), strides=3)(x)  
aux1 = layers.Conv2D(128, 1, padding='same', activation='relu')(aux1)  
aux1 = layers.Flatten()(aux1)  
aux1 = layers.Dense(1024, activation='relu')(aux1)  
aux1 = layers.Dropout(0.7)(aux1)  
aux1 = layers.Dense(10, activation='softmax')(aux1)
```

```
x = inception(x,  
    filters_1x1=160,  
    filters_3x3_reduce=112,  
    filters_3x3=224,  
    filters_5x5_reduce=24,  
    filters_5x5=64,  
    filters_pool=64)
```

```
x = inception(x,  
    filters_1x1=128,  
    filters_3x3_reduce=128,  
    filters_3x3=256,  
    filters_5x5_reduce=24,  
    filters_5x5=64,  
    filters_pool=64)
```

```
x = inception(x,  
    filters_1x1=112,  
    filters_3x3_reduce=144,  
    filters_3x3=288,  
    filters_5x5_reduce=32,  
    filters_5x5=64,  
    filters_pool=64)
```

```
aux2 = layers.AveragePooling2D((5, 5), strides=3)(x)  
aux2 = layers.Conv2D(128, 1, padding='same', activation='relu')(aux2)  
aux2 = layers.Flatten()(aux2)  
aux2 = layers.Dense(1024, activation='relu')(aux2)  
aux2 = layers.Dropout(0.7)(aux2)  
aux2 = layers.Dense(10, activation='softmax')(aux2)
```

```
x = inception(x,  
    filters_1x1=256,  
    filters_3x3_reduce=160,  
    filters_3x3=320,  
    filters_5x5_reduce=32,  
    filters_5x5=128,  
    filters_pool=128)
```

```
x = layers.MaxPooling2D(3, strides=2)(x)
```

```
x = inception(x,  
    filters_1x1=256,  
    filters_3x3_reduce=160,  
    filters_3x3=320,  
    filters_5x5_reduce=32,  
    filters_5x5=128,
```

```
filters_pool=128)
```

```
x = inception(x,  
              filters_1x1=384,  
              filters_3x3_reduce=192,  
              filters_3x3=384,  
              filters_5x5_reduce=48,  
              filters_5x5=128,  
              filters_pool=128)
```

```
x = layers.GlobalAveragePooling2D()(x)
```

```
x = layers.Dropout(0.4)(x)  
out = layers.Dense(10, activation='softmax')(x)
```

```
model = Model(inputs = inp, outputs = [out, aux1, aux2])
```

```
model.compile(optimizer='adam', loss=[losses.sparse_categorical_crossentropy,  
losses.sparse_categorical_crossentropy, losses.sparse_categorical_crossentropy],  
loss_weights=[1, 0.3, 0.3], metrics=['accuracy'])
```

```
history = model.fit(X_train, [y_train, y_train, y_train], validation_data=(X_test,  
[y_test, y_test, y_test]), batch_size=64, epochs=30)
```

```
fig, axs = plt.subplots(figsize=(12,8))
```

```
axs.plot(history.history['loss'])  
axs.plot(history.history['val_loss'])  
axs.title.set_text('Training Loss vs Validation Loss')  
axs.set_xlabel('Epochs')  
axs.set_ylabel('Loss')
```



```
axs.legend(['Train','Val'])
```

```
plt.show()
```

```
model.evaluate(X_test, y_test)
```

```
# EmoDB Dataset
```

```
!unzip "/content/drive/MyDrive/EmoDB.zip"
```

```
import librosa
```

```
import numpy as np
```

```
input_length = 16000*5
```

```
batch_size = 32
```

```
n_mels = 320
```

```
def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,
```

```
#log_specgram
```

```
    step_size=10, eps=1e-10):
```

```
    mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,  
n_mels= n_mels)
```

```
    mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40
```

```
    return mel_db.T
```

```
def load_audio_file(file_path, input_length=input_length):
```

```
data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000
if len(data)>input_length:
    max_offset = len(data)-input_length

    offset = np.random.randint(max_offset)

    data = data[offset:(input_length+offset)]

else:
    if input_length > len(data):
        max_offset = input_length - len(data)

        offset = np.random.randint(max_offset)
    else:
        offset = 0
    data = np.pad(data, (offset, input_length - len(data) - offset), "constant")

data = preprocess_audio_mel_T(data)
return data
```

Preprocessing the dataset

```
import os
from scipy.io import wavfile
import librosa
import matplotlib.pyplot as plt
import numpy as np
import cv2
```

```
directory = "/content/wav/"
```

```
classes = ["W" ,"L" ,"E" ,"A" , "F" ,"T" ,"N" ]
```

```
X = list()
```

```
y = list()
```

```
for filename in os.listdir(directory):
```

```
    filePath = os.path.join(directory, filename)
```

```
    a = load_audio_file(file_path=filePath)
```

```
    data = cv2.merge([a,a,a])
```

```
    if(filename[5:6] in classes):
```

```
        X.append(data)
```

```
        y.append(classes.index(filename[5:6]))
```

```
X = np.asarray(X, dtype=np.float32)
```

```
y = np.asarray(y, dtype=np.float32)
```

```
import pandas as pd
```

```
import numpy as np
```

```
import matplotlib.pyplot as plt
```

```
import tensorflow as tf
```

```
# dataset preparation
```

```
from tensorflow.keras import datasets, layers, models
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.4, train_size= 0.6  
, random_state=10)
```

```
def inception(x,
```

```
    filters_1x1,
```

```
    filters_3x3_reduce,
```

```
    filters_3x3,
```

```
    filters_5x5_reduce,
```

```
    filters_5x5,
```

```

        filters_pool):
    path1 = layers.Conv2D(filters_1x1, (1, 1), padding='same', activation='relu')(x)

    path2 = layers.Conv2D(filters_3x3_reduce, (1, 1), padding='same',
activation='relu')(x)
    path2 = layers.Conv2D(filters_3x3, (1, 1), padding='same',
activation='relu')(path2)

    path3 = layers.Conv2D(filters_5x5_reduce, (1, 1), padding='same',
activation='relu')(x)
    path3 = layers.Conv2D(filters_5x5, (1, 1), padding='same',
activation='relu')(path3)

    path4 = layers.MaxPool2D((3, 3), strides=(1, 1), padding='same')(x)
    path4 = layers.Conv2D(filters_pool, (1, 1), padding='same',
activation='relu')(path4)

    return tf.concat([path1, path2, path3, path4], axis=3)

inp = layers.Input(shape=(157, 320, 3))
input_tensor = layers.experimental.preprocessing.Resizing(224, 224,
interpolation="bilinear", input_shape=X_train.shape[1:])(inp)

x = layers.Conv2D(64, 7, strides=2, padding='same',
activation='relu')(input_tensor)
x = layers.MaxPooling2D(3, strides=2)(x)

x = layers.Conv2D(64, 1, strides=1, padding='same', activation='relu')(x)
x = layers.Conv2D(192, 3, strides=1, padding='same', activation='relu')(x)

x = layers.MaxPooling2D(3, strides=2)(x)

```

```
x = inception(x,  
    filters_1x1=64,  
    filters_3x3_reduce=96,  
    filters_3x3=128,  
    filters_5x5_reduce=16,  
    filters_5x5=32,  
    filters_pool=32)
```

```
x = inception(x,  
    filters_1x1=128,  
    filters_3x3_reduce=128,  
    filters_3x3=192,  
    filters_5x5_reduce=32,  
    filters_5x5=96,  
    filters_pool=64)
```

```
x = layers.MaxPooling2D(3, strides=2)(x)
```

```
x = inception(x,  
    filters_1x1=192,  
    filters_3x3_reduce=96,  
    filters_3x3=208,  
    filters_5x5_reduce=16,  
    filters_5x5=48,  
    filters_pool=64)
```

```
aux1 = layers.AveragePooling2D((5, 5), strides=3)(x)  
aux1 = layers.Conv2D(128, 1, padding='same', activation='relu')(aux1)  
aux1 = layers.Flatten()(aux1)  
aux1 = layers.Dense(1024, activation='relu')(aux1)  
aux1 = layers.Dropout(0.7)(aux1)
```

```
aux1 = layers.Dense(10, activation='softmax')(aux1)
```

```
x = inception(x,  
    filters_1x1=160,  
    filters_3x3_reduce=112,  
    filters_3x3=224,  
    filters_5x5_reduce=24,  
    filters_5x5=64,  
    filters_pool=64)
```

```
x = inception(x,  
    filters_1x1=128,  
    filters_3x3_reduce=128,  
    filters_3x3=256,  
    filters_5x5_reduce=24,  
    filters_5x5=64,  
    filters_pool=64)
```

```
x = inception(x,  
    filters_1x1=112,  
    filters_3x3_reduce=144,  
    filters_3x3=288,  
    filters_5x5_reduce=32,  
    filters_5x5=64,  
    filters_pool=64)
```

```
aux2 = layers.AveragePooling2D((5, 5), strides=3)(x)  
aux2 = layers.Conv2D(128, 1, padding='same', activation='relu')(aux2)  
aux2 = layers.Flatten()(aux2)  
aux2 = layers.Dense(1024, activation='relu')(aux2)  
aux2 = layers.Dropout(0.7)(aux2)  
aux2 = layers.Dense(10, activation='softmax')(aux2)
```

```
x = inception(x,  
              filters_1x1=256,  
              filters_3x3_reduce=160,  
              filters_3x3=320,  
              filters_5x5_reduce=32,  
              filters_5x5=128,  
              filters_pool=128)
```

```
x = layers.MaxPooling2D(3, strides=2)(x)
```

```
x = inception(x,  
              filters_1x1=256,  
              filters_3x3_reduce=160,  
              filters_3x3=320,  
              filters_5x5_reduce=32,  
              filters_5x5=128,  
              filters_pool=128)
```

```
x = inception(x,  
              filters_1x1=384,  
              filters_3x3_reduce=192,  
              filters_3x3=384,  
              filters_5x5_reduce=48,  
              filters_5x5=128,  
              filters_pool=128)
```

```
x = layers.GlobalAveragePooling2D()(x)
```

```
x = layers.Dropout(0.4)(x)
```

```
out = layers.Dense(10, activation='softmax')(x)
```

```
model = Model(inputs = inp, outputs = [out, aux1, aux2])
```

```
model.compile(optimizer='adam', loss=[losses.sparse_categorical_crossentropy,  
losses.sparse_categorical_crossentropy, losses.sparse_categorical_crossentropy],  
loss_weights=[1, 0.3, 0.3], metrics=['accuracy'])
```

```
history = model.fit(X_train, [y_train, y_train, y_train], validation_data=(X_test,  
[y_test, y_test, y_test]), batch_size=64, epochs=30)
```

```
fig, axs = plt.subplots(figsize=(12,8))
```

```
axs.plot(history.history['loss'])
```

```
axs.plot(history.history['val_loss'])
```

```
axs.title.set_text('Training Loss vs Validation Loss')
```

```
axs.set_xlabel('Epochs')
```

```
axs.set_ylabel('Loss')
```

```
axs.legend(['Train','Val'])
```

```
plt.show()
```

```
model.evaluate(X_test, y_test)
```

```
from google.colab import drive
```

```
drive.mount('/content/drive')
```

```
# Import necessary packages
```

```
import argparse
```

```
# Import necessary components to build LeNet
```

```
from keras.models import Sequential
```

```
from keras.layers.core import Dense, Dropout, Activation, Flatten
```

```
from keras.layers.convolutional import Conv2D, MaxPooling2D, ZeroPadding2D
```



```
from keras.layers import BatchNormalization
from keras.regularizers import l2

import tensorflow as tf
from tensorflow import keras
import matplotlib.pyplot as plt
%matplotlib inline
import numpy as np
import skimage.transform

def alexnet_model(img_shape=(224, 224, 3), n_classes=10, l2_reg=0.,
                  weights=None):

    # Initialize model
    alexnet = Sequential()

    # Layer 1
    alexnet.add(Conv2D(30, (11, 11), input_shape=img_shape,
                      padding='same', kernel_regularizer=l2(l2_reg)))
    alexnet.add(BatchNormalization())
    alexnet.add(Activation('relu'))
    alexnet.add(MaxPooling2D(pool_size=(2, 2)))

    # Layer 2
    alexnet.add(Conv2D(30, (5, 5), padding='same'))
    alexnet.add(BatchNormalization())
    alexnet.add(Activation('relu'))
    alexnet.add(MaxPooling2D(pool_size=(2, 2)))

    # Layer 3
    alexnet.add(ZeroPadding2D((1, 1)))
    alexnet.add(Conv2D(30, (3, 3), padding='same'))
```

```
alexnet.add(BatchNormalization())
alexnet.add(Activation('relu'))
alexnet.add(MaxPooling2D(pool_size=(2, 2)))
```

Layer 4

```
alexnet.add(ZeroPadding2D((1, 1)))
alexnet.add(Conv2D(30, (3, 3), padding='same'))
alexnet.add(BatchNormalization())
alexnet.add(Activation('relu'))
```

Layer 5

```
alexnet.add(ZeroPadding2D((1, 1)))
alexnet.add(Conv2D(30, (3, 3), padding='same'))
alexnet.add(BatchNormalization())
alexnet.add(Activation('relu'))
alexnet.add(MaxPooling2D(pool_size=(2, 2)))
```

Layer 6

```
alexnet.add(Flatten())
alexnet.add(Dense(30))
alexnet.add(BatchNormalization())
alexnet.add(Activation('relu'))
alexnet.add(Dropout(0.5))
```

Layer 7

```
alexnet.add(Dense(30))
alexnet.add(BatchNormalization())
alexnet.add(Activation('relu'))
alexnet.add(Dropout(0.5))
```

Layer 8

```
alexnet.add(Dense(n_classes))
```

```
alexnet.add(BatchNormalization())  
alexnet.add(Activation('softmax'))
```

```
if weights is not None:  
    alexnet.load_weights(weights)
```

```
return alexnet
```

```
def parse_args():
```

```
    """
```

```
    Parse command line arguments.
```

```
    Parameters:
```

```
        None
```

```
    Returns:
```

```
        parser arguments
```

```
    """
```

```
    parser = argparse.ArgumentParser(description='AlexNet model')
```

```
    optional = parser._action_groups.pop()
```

```
    required = parser.add_argument_group('required arguments')
```

```
    optional.add_argument('--print_model',
```

```
        dest='print_model',
```

```
        help='Print AlexNet model',
```

```
        action='store_true')
```

```
    parser._action_groups.append(optional)
```

```
    return parser.parse_args()
```

```
def load_preprocess_training_batch(X_train):
```

```
    new = []
```

```
    for item in X_train:
```

```
        tmpFeature = skimage.transform.resize(item, (224, 224), mode='constant')
```

```
new.append(tmpFeature)
```

```
return new
```

```
# CIFAR 10 DATASET
```

```
# Command line parameters
```

```
# args = parse_args()
```

```
# Create AlexNet model
```

```
model = alexnet_model()
```

```
# Print
```

```
# if args.print_model:
```

```
#     model.summary()
```

```
(X_train, y_train) , (X_test, y_test) = keras.datasets.cifar10.load_data()
```

```
X_train = X_train[0:500]
```

```
y_train = y_train[0:500]
```

```
X_test = X_test[0:200]
```

```
y_test = y_test[0:200]
```

```
X_train_resized = load_preprocess_training_batch(X_train)
```

```
X_test_resized = load_preprocess_training_batch(X_test)
```

```
X_train_resized = np.array(X_train_resized)
```

```
X_test_resized = np.array(X_test_resized)
```

```
X_train_resized = X_train_resized / 255
```

```
X_test_resized = X_test_resized / 255
```

```
model.compile(optimizer='SGD',  
              loss='sparse_categorical_crossentropy',  
              metrics=['accuracy'])
```

```
history = model.fit(X_train_resized, y_train, epochs=5)
```

```
model.evaluate(X_test_resized, y_test)
```

```
# NMIST Dataset
```

```
(X_train, y_train) , (X_test, y_test) = keras.datasets.mnist.load_data()
```

```
X_train = X_train[0:2000]
```

```
y_train = y_train[0:2000]
```

```
X_test = X_test[0:2000]
```

```
y_test = y_test[0:2000]
```

```
X_train_resized = load_preprocess_training_batch(X_train)
```

```
X_test_resized = load_preprocess_training_batch(X_test)
```

```
X_train_resized = np.array(X_train_resized)
```

```
X_test_resized = np.array(X_test_resized)
```

```
X_train_resized = X_train_resized / 255.0
```

```
X_test_resized = X_test_resized / 255.0
```

```
import cv2
```

```
X_train_new = list()
```

```
for i in range(len(X_train_resized)):
```

```
    g = X_train_resized[i]
```

```
X_train_new.append(cv2.merge([g,g,g]))

X_train_new = np.asarray(X_train_new,dtype=np.float32)

X_test_new = list()

for i in range(len(X_test_resized)):
    g = X_test_resized[i]
    X_test_new.append(cv2.merge([g,g,g]))

X_test_new = np.asarray(X_test_new,dtype=np.float32)

model = alexnet_model()

model.compile(optimizer='SGD',
               loss='sparse_categorical_crossentropy',
               metrics=['accuracy'])

history = model.fit(X_train_new, y_train, epochs=5)

model.evaluate(X_test_new, y_test)

# SAVEE Dataset

!unzip "/content/drive/MyDrive/SaveeDataset.zip"

import librosa
import numpy as np

input_length = 16000*5

batch_size = 32
```

```
n_mels = 320
```

```
def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,  
#log_specgram  
    step_size=10, eps=1e-10):
```

```
    mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,  
n_mels= n_mels)
```

```
    mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40
```

```
    return mel_db.T
```

```
def load_audio_file(file_path, input_length=input_length):
```

```
    data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000
```

```
    if len(data)>input_length:
```

```
        max_offset = len(data)-input_length
```

```
        offset = np.random.randint(max_offset)
```

```
        data = data[offset:(input_length+offset)]
```

```
    else:
```

```
        if input_length > len(data):
```

```
            max_offset = input_length - len(data)
```

```
            offset = np.random.randint(max_offset)
```

```
            else:
```

```
                offset = 0
```

```
                data = np.pad(data, (offset, input_length - len(data) - offset), "constant")
```

```
data = preprocess_audio_mel_T(data)
return data

# Preprocessing the dataset
import os
from scipy.io import wavfile
import librosa
import matplotlib.pyplot as plt
import numpy as np
import cv2

rootDirectory = "/content/AudioData/"
personNames = ["DC","JE","JK","KL"]

classes = ["a" , "d" , "f" , "h" , "n" , "sa" , "su" ]

X = list()
y = list()

for person in personNames:
    directory = os.path.join(rootDirectory,person)
    for filename in os.listdir(directory):
        filePath = os.path.join(directory, filename)
        a = load_audio_file(file_path=filePath)
        data = cv2.merge([a,a,a])
        if(filename[0:1] in classes):
            X.append(data)
            y.append(classes.index(filename[0:1]))
        elif(filename[0:2] in classes):
            X.append(data)
            y.append(classes.index(filename[0:2]))
```



```
X = np.asarray(X, dtype=np.float32)
y = np.asarray(y, dtype=np.float32)

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import tensorflow as tf

# dataset preparation

from tensorflow.keras import datasets, layers, models
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.5, train_size= 0.5
, random_state=10)

X_train_resized = load_preprocess_training_batch(X_train)
X_test_resized = load_preprocess_training_batch(X_test)

X_train_resized = np.array(X_train_resized)
X_test_resized = np.array(X_test_resized)

model = alexnet_model()

model.compile(optimizer='SGD',
              loss='sparse_categorical_crossentropy',
              metrics=['accuracy'])

history = model.fit(X_train_resized, y_train, epochs=10)

model.evaluate(X_test_resized, y_test)

# EmoDB Database
```

```
!unzip "/content/drive/MyDrive/EmoDB.zip"
```

```
import librosa  
import numpy as np
```

```
input_length = 16000*5
```

```
batch_size = 32
```

```
n_mels = 320
```

```
def preprocess_audio_mel_T(audio, sample_rate=16000, window_size=20,  
#log_specgram  
    step_size=10, eps=1e-10):
```

```
    mel_spec = librosa.feature.melspectrogram(y=audio, sr=sample_rate,  
n_mels= n_mels)
```

```
    mel_db = (librosa.power_to_db(mel_spec, ref=np.max) + 40)/40
```

```
    return mel_db.T
```

```
def load_audio_file(file_path, input_length=input_length):
```

```
    data = librosa.core.load(file_path, sr=16000)[0] #, sr=16000
```

```
    if len(data)>input_length:
```

```
        max_offset = len(data)-input_length
```

```
        offset = np.random.randint(max_offset)
```

```
        data = data[offset:(input_length+offset)]
```

```
else:
    if input_length > len(data):
        max_offset = input_length - len(data)

        offset = np.random.randint(max_offset)
    else:
        offset = 0
    data = np.pad(data, (offset, input_length - len(data) - offset), "constant")

data = preprocess_audio_mel_T(data)
return data
```

Preprocessing the dataset

```
import os
from scipy.io import wavfile
import librosa
import matplotlib.pyplot as plt
import numpy as np
import cv2
```

```
directory = "/content/wav/"
```

```
classes = ["W" , "L" , "E" , "A" , "F" , "T" , "N" ]
```

```
X = list()
```

```
y = list()
```

```
for filename in os.listdir(directory):
    filePath = os.path.join(directory, filename)
    a = load_audio_file(file_path=filePath)
    data = cv2.merge([a,a,a])
    if(filename[5:6] in classes):
```

```
X.append(data)  
y.append(classes.index(filename[5:6]))
```

```
X = np.asarray(X, dtype=np.float32)  
y = np.asarray(y, dtype=np.float32)
```

```
import pandas as pd  
import numpy as np  
import matplotlib.pyplot as plt  
import tensorflow as tf
```

```
# dataset preparation
```

```
from tensorflow.keras import datasets, layers, models  
from sklearn.model_selection import train_test_split  
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.4, train_size= 0.6  
, random_state=10)
```

```
X_train_resized = load_preprocess_training_batch(X_train)  
X_test_resized = load_preprocess_training_batch(X_test)
```

```
X_train_resized = np.array(X_train_resized)  
X_test_resized = np.array(X_test_resized)
```

```
model = alexnet_model()
```

```
model.compile(optimizer='SGD',  
              loss='sparse_categorical_crossentropy',  
              metrics=['accuracy'])
```

```
history = model.fit(X_train_resized, y_train, epochs=10)
```

```
model.evaluate(X_test_resized, y_test)
```